

This Page Is Inserted by IFW Operations  
and is not a part of the Official Record

## **BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

**IMAGES ARE BEST AVAILABLE COPY.**

**As rescanning documents *will not* correct images,  
please do not report the images to the  
Image Problem Mailbox.**

(19) World Intellectual Property Organization  
International Bureau



(43) International Publication Date  
7 February 2002 (07.02.2002)

PCT

(10) International Publication Number  
**WO 02/09755 A2**

- (51) International Patent Classification<sup>7</sup>: **A61K 39/395**
- (21) International Application Number: **PCT/US01/23691**
- (22) International Filing Date: **27 July 2001 (27.07.2001)**
- (25) Filing Language: **English**
- (26) Publication Language: **English**
- (30) Priority Data:  
60/221,256      27 July 2000 (27.07.2000)      **US**
- (71) Applicant (*for all designated States except US*): **GENENTECH, INC.** [US/US]; 1 DNA Way, South San Francisco, CA 94080 (US).
- (72) Inventors; and
- (75) Inventors/Applicants (*for US only*): **ESCANDON, Enrique** [MX/US]; 10 Garden Ct. Apt. #2, Belmont, CA 94002 (US). **FOX, Judith, A.** [US/US]; 1531 Golden Gate Avenue, San Francisco, CA 94115 (US). **KELLEY, Sean, K.** [US/US]; 3915 Edison Street, San Mateo, CA 94403 (US). **XIANG, Hong** [US/US]; 558 Driscoll Place, Palo Alto, CA 94306 (US).
- (74) Agent: **MARSCHANG, Diane, L.**; Genentech, Inc., MS 49, 1 DNA Way, South San Francisco, CA 94080 (US).
- (81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).
- Published:  
— *without international search report and to be republished upon receipt of that report*
- For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.*



**WO 02/09755 A2**

(54) Title: **APO-2L RECEPTOR AGONIST AND CPT-11 SYNERGISM**

(57) Abstract: Methods of using effective amounts of Apo-2L receptor agonists and CPT-11 to induce apoptosis and suppress growth of cancer cells are provided.

Apo-2L RECEPTOR AGONIST and CPT-11 SYNERGISM

5

FIELD OF THE INVENTION

This invention relates generally to methods of inducing apoptosis in mammalian cells. In particular, it pertains to the use of Apo-2L receptor agonists and CPT-11 to synergistically induce apoptosis in mammalian cells. Various Apo-2L receptor agonists contemplated by the invention include the ligand known as Apo-2 ligand or TRAIL, as well as agonist antibodies directed to one or more Apo-2L receptors.

BACKGROUND OF THE INVENTION

Various molecules, such as tumor necrosis factor- $\alpha$  ("TNF- $\alpha$ "), tumor necrosis factor- $\beta$  ("TNF- $\beta$ " or "lymphotoxin- $\alpha$ "), lymphotoxin- $\beta$  ("LT- $\beta$ "), CD30 ligand, CD27 ligand, CD40 ligand, OX-40 ligand, 4-1BB ligand, Apo-1 ligand (also referred to as Fas ligand or CD95 ligand), Apo-2 ligand (also referred to as TRAIL), Apo-3 ligand (also referred to as TWEAK), osteoprotegerin (OPG), APRIL, RANK ligand (also referred to as TRANCE), and TALL-1 (also referred to as BlyS, BAFF or THANK) have been identified as members of the tumor necrosis factor ("TNF") family of cytokines [See, e.g., Gruss and Dower, Blood, 85:3378-3404 (1995); Pitti et al., J. Biol. Chem., 271:12687-12690 (1996); Wiley et al., Immunity, 3:673-682 (1995); Browning et al., Cell, 72:847-856 (1993); Armitage et al., Nature, 357:80-82 (1992), WO 97/01633 published January 16, 1997; WO 97/25428 published July 17, 1997; Marsters et al., Curr. Biol., 8:525-528 (1998); Simonet et al., Cell, 89:309-319 (1997); Chicheportiche et al., Biol. Chem., 272:32401-32410 (1997); Hahne et al., J. Exp. Med., 188:1185-1190 (1998); WO98/28426 published July 2, 1998; WO98/46751 published October 22, 1998; WO/98/18921 published May 7, 1998; Moore et al., Science, 285:260-263 (1999); Shu et al., J. Leukocyte Biol., 65:680 (1999); Schneider et al., J. Exp. Med., 189:1747-1756 (1999); Mukhopadhyay et al., J. Biol. Chem., 274:15978-15981 (1999)]. Among these molecules, TNF- $\alpha$ , TNF- $\beta$ , CD30 ligand, 4-1BB ligand, Apo-1 ligand, Apo-2 ligand (Apo2L/TRAIL) and Apo-3 ligand (TWEAK) have been reported to be involved in apoptotic cell death. Both TNF- $\alpha$  and TNF- $\beta$  have been reported to induce apoptotic death in susceptible tumor cells [Schmid et al., Proc. Natl. Acad. Sci., 83:1881 (1986); Dealtry et al., Eur. J. Immunol., 17:689 (1987)].

Recently, additional molecules believed to be members of the TNF cytokine family were identified and reported to be involved in apoptosis. For instance, in Pitti et al., J. Biol. Chem., 271:12687-12690 (1996), a molecule referred to as Apo-2 ligand is described. See also, WO 97/25428 published July 17, 1997. The full length human Apo-2 ligand is reported to be a 281 amino acid polypeptide that induces apoptosis in various mammalian cells. Other investigators have described related polypeptides

referred to as TRAIL [Wiley et al., Immunity, 3:673-682 (1995); WO 97/01633 published January 16, 1997] and AGP-1 [WO 97/46686 published December 11, 1997].

5 Various molecules in the TNF family also have purported role(s) in the function or development of the immune system [Gruss et al., Blood, 85:3378 (1995)]. Zheng et al. have reported that TNF- $\alpha$  is involved in post-stimulation apoptosis of CD8-positive T cells [Zheng et al., Nature, 377:348-351 (1995)]. Other investigators have reported that CD30 ligand may be involved in deletion of self-reactive T cells in the thymus  
10 [Amakawa et al., Cold Spring Harbor Laboratory Symposium on Programmed Cell Death, Abstr. No. 10, (1995)]. CD40 ligand activates many functions of B cells, including proliferation, immunoglobulin secretion, and survival [Renshaw et al., J. Exp. Med., 180:1889 (1994)]. Another recently identified TNF family cytokine, TALL-1 (BlyS), has been  
15 reported, under certain conditions, to induce B cell proliferation and immunoglobulin secretion. [Moore et al., supra; Schneider et al., supra; Mackay et al., J. Exp. Med., 190:1697 (1999)].

Mutations in the mouse Fas/Apo-1 receptor or ligand genes (called *lpr* and *gld*, respectively) have been associated with some autoimmune  
20 disorders, indicating that Apo-1 ligand may play a role in regulating the clonal deletion of self-reactive lymphocytes in the periphery [Krammer et al., Curr. Op. Immunol., 6:279-289 (1994); Nagata et al., Science, 267:1449-1456 (1995)]. Apo-1 ligand is also reported to induce post-stimulation apoptosis in CD4-positive T lymphocytes and in B lymphocytes,  
25 and may be involved in the elimination of activated lymphocytes when their function is no longer needed [Krammer et al., supra; Nagata et al., supra]. Agonist mouse monoclonal antibodies specifically binding to the Apo-1 receptor have been reported to exhibit cell killing activity that is comparable to or similar to that of TNF- $\alpha$  [Yonehara et al., J. Exp.  
30 Med., 169:1747-1756 (1989)].

Induction of various cellular responses mediated by such TNF family cytokines is believed to be initiated by their binding to specific cell receptors. Previously, two distinct TNF receptors of approximately 55-kDa (TNFR1) and 75-kDa (TNFR2) were identified [Hohman et al., J. Biol.  
35 Chem., 264:14927-14934 (1989); Brockhaus et al., Proc. Natl. Acad. Sci., 87:3127-3131 (1990); EP 417,563, published March 20, 1991; Loetscher et al., Cell, 61:351 (1990); Schall et al., Cell, 61:361 (1990); Smith et al., Science, 248:1019-1023 (1990); Lewis et al., Proc. Natl. Acad. Sci., 88:2830-2834 (1991); Goodwin et al., Mol. Cell. Biol., 11:3020-3026  
40 (1991)]. Those TNFRs were found to share the typical structure of cell surface receptors including extracellular, transmembrane and intracellular regions. The extracellular portions of both receptors were found naturally also as soluble TNF-binding proteins [Nophar, Y. et al., EMBO J., 9:3269 (1990); and Kohno, T. et al., Proc. Natl. Acad. Sci.



U.S.A., 87:8331 (1990); Hale et al., J. Cell. Biochem. Supplement 15F, 1991, p. 113 (P424)].

The extracellular portion of type 1 and type 2 TNFRs (TNFR1 and TNFR2) contains a repetitive amino acid sequence pattern of four  
5 cysteine-rich domains (CRDs) designated 1 through 4, starting from the NH<sub>2</sub>-terminus. [Schall et al., supra; Loetscher et al., supra; Smith et al., supra; Nophar et al., supra; Kohno et al., supra; Banner et al., Cell, 73:431-435 (1993)]. A similar repetitive pattern of CRDs exists in several other cell-surface proteins, including the p75 nerve growth  
10 factor receptor (NGFR) [Johnson et al., Cell, 47:545 (1986); Radeke et al., Nature, 325:593 (1987)], the B cell antigen CD40 [Stamenkovic et al., EMBO J., 8:1403 (1989)], the T cell antigen OX40 [Mallet et al., EMBO J., 9:1063 (1990)] and the Fas antigen [Yonehara et al., supra and Itoh et al., Cell, 66:233-243 (1991)]. CRDs are also found in the  
15 soluble TNFR (sTNFR)-like T2 proteins of the Shope and myxoma poxviruses [Upton et al., Virology, 160:20-29 (1987); Smith et al., Biochem. Biophys. Res. Commun., 176:335 (1991); Upton et al., Virology, 184:370 (1991)]. Optimal alignment of these sequences indicates that the positions of the cysteine residues are well conserved. These receptors  
20 are sometimes collectively referred to as members of the TNF/NGF receptor superfamily.

The TNF family ligands identified to date, with the exception of lymphotoxin- $\alpha$ , are type II transmembrane proteins, whose C-terminus is extracellular. In contrast, most receptors in the TNF receptor (TNFR)  
25 family identified to date are type I transmembrane proteins. In both the TNF ligand and receptor families, however, homology identified between family members has been found mainly in the extracellular domain ("ECD"). Several of the TNF family cytokines, including TNF- $\alpha$ , Apo-1 ligand and CD40 ligand, are cleaved proteolytically at the cell surface; the  
30 resulting protein in each case typically forms a homotrimeric molecule that functions as a soluble cytokine. TNF receptor family proteins are also usually cleaved proteolytically to release soluble receptor ECDs that can function as inhibitors of the cognate cytokines.

More recently, other members of the TNFR family have been  
35 identified. In von Bulow et al., Science, 278:138-141 (1997), investigators describe a plasma membrane receptor referred to as Transmembrane Activator and CAML-Interactor or "TACI". The TACI receptor is reported to contain a cysteine-rich motif characteristic of the TNFR family. In an in vitro assay, cross linking of TACI on the surface of  
40 transfected Jurkat cells with TACI-specific antibodies led to activation of NF-KB. [see also, WO 98/39361 published September 18, 1998].

Laabi et al., EMBO J., 11:3897-3904 (1992) reported identifying a new gene called "BCM" whose expression was found to coincide with B cell terminal maturation. The open reading frame of the BCM normal cDNA  
45 predicted a 184 amino acid long polypeptide with a single transmembrane

domain. These investigators later termed this gene "BCMA." [Laabi et al., Nucleic Acids Res., 22:1147-1154 (1994)]. BCMA mRNA expression was reported to be absent in human malignant B cell lines which represent the pro-B lymphocyte stage, and thus, is believed to be linked to the stage of differentiation of lymphocytes [Gras et al., Int. Immunology, 7:1093-1106 (1995)]. In Madry et al., Int. Immunology, 10:1693-1702 (1998), the cloning of murine BCMA cDNA was described. The murine BCMA cDNA is reported to encode a 185 amino acid long polypeptide having 62% identity to the human BCMA polypeptide. Alignment of the murine and human BCMA protein sequences revealed a conserved motif of six cysteines in the N-terminal region, suggesting that the BCMA protein belongs to the TNFR superfamily [Madry et al., supra].

In Marsters et al., Curr. Biol., 6:750 (1996), investigators describe a full length native sequence human polypeptide, called Apo-3, which exhibits similarity to the TNFR family in its extracellular cysteine-rich repeats and resembles TNFR1 and CD95 in that it contains a cytoplasmic death domain sequence [see also Marsters et al., Curr. Biol., 6:1669 (1996)]. Apo-3 has also been referred to by other investigators as DR3, wsl-1, TRAMP, and LARD [Chinnaiyan et al., Science, 274:990 (1996); Kitson et al., Nature, 384:372 (1996); Bodmer et al., Immunity, 6:79 (1997); Screaton et al., Proc. Natl. Acad. Sci., 94:4615-4619 (1997)].

Pan et al. have disclosed another TNF receptor family member referred to as "DR4" [Pan et al., Science, 276:111-113 (1997); see also WO98/32856 published July 30, 1998]. The DR4 was reported to contain a cytoplasmic death domain capable of engaging the cell suicide apparatus. Pan et al. disclose that DR4 is believed to be a receptor for the ligand known as Apo2L/TRAIL.

In Sheridan et al., Science, 277:818-821 (1997) and Pan et al., Science, 277:815-818 (1997), another molecule believed to be a receptor for Apo2L/TRAIL is described [see also, WO98/51793 published November 19, 1998; WO98/41629 published September 24, 1998]. That molecule is referred to as DR5 (it has also been alternatively referred to as Apo-2; TRAIL-R, TR6, Tango-63, hAP08, TRICK2 or KILLER [Screaton et al., Curr. Biol., 7:693-696 (1997); Walczak et al., EMBO J., 16:5386-5387 (1997); Wu et al., Nature Genetics, 17:141-143 (1997); WO98/35986 published August 20, 1998; EP870,827 published October 14, 1998; WO98/46643 published October 22, 1998; WO99/02653 published January 21, 1999; WO99/09165 published February 25, 1999; WO99/11791 published March 11, 1999]. Like DR4, DR5 is reported to contain a cytoplasmic death domain and be capable of signaling apoptosis. The crystal structure of the complex formed between Apo-2L/TRAIL and DR5 is described in Hymowitz et al., Molecular Cell, 4:563-571 (1999).

Yet another death domain-containing receptor, DR6, was recently identified [Pan et al., FEBS Letters, 431:351-356 (1998)]. Aside from

containing four putative extracellular cysteine rich domains and a cytoplasmic death domain, DR6 is believed to contain a putative leucine-zipper sequence that overlaps with a proline-rich motif in the cytoplasmic region. The proline-rich motif resembles sequences that bind  
5 to src-homology-3 domains, which are found in many intracellular signal-transducing molecules.

A further group of recently identified receptors are referred to as "decoy receptors," which are believed to function as inhibitors, rather than transducers of signaling. This group includes DCR1 (also referred  
10 to as TRID, LIT or TRAIL-R3) [Pan et al., Science, 276:111-113 (1997); Sheridan et al., Science, 277:818-821 (1997); McFarlane et al., J. Biol. Chem., 272:25417-25420 (1997); Schneider et al., FEBS Letters, 416:329-334 (1997); Degli-Esposti et al., J. Exp. Med., 186:1165-1170 (1997); and Mongkolsapaya et al., J. Immunol., 160:3-6 (1998)] and DCR2 (also called  
15 TRUNDD or TRAIL-R4) [Marsters et al., Curr. Biol., 7:1003-1006 (1997); Pan et al., FEBS Letters, 424:41-45 (1998); Degli-Esposti et al., Immunity, 7:813-820 (1997)], both cell surface molecules, as well as OPG [Simonet et al., supra; Emery et al., infra] and DCR3 [Pitti et al., Nature, 396:699-703 (1998)], both of which are secreted, soluble  
20 proteins.

Additional newly identified members of the TNFR family include CAR1, HVEM, GITR, ZTNFR-5, NTR-1, and TNFL1 [Brojatsch et al., Cell, 87:845-855 (1996); Montgomery et al., Cell, 87:427-436 (1996); Marsters et al., J. Biol. Chem., 272:14029-14032 (1997); Nocentini et al., Proc.  
25 Natl. Acad. Sci. USA 94:6216-6221 (1997); Emery et al., J. Biol. Chem., 273:14363-14367 (1998); WO99/04001 published January 28, 1999; WO99/07738 published February 18, 1999; WO99/33980 published July 8, 1999].

As reviewed recently by Tewari et al., TNFR1, TNFR2 and CD40 modulate the expression of proinflammatory and costimulatory cytokines,  
30 cytokine receptors, and cell adhesion molecules through activation of the transcription factor, NF-KB [Tewari et al., Curr. Op. Genet. Develop., 6:39-44 (1996)]. NF-KB is the prototype of a family of dimeric transcription factors whose subunits contain conserved Rel regions [Verma et al., Genes Develop., 9:2723-2735 (1996); Baldwin, Ann. Rev. Immunol.,  
35 14:649-681 (1996)]. In its latent form, NF-KB is complexed with members of the IKB inhibitor family; upon inactivation of the IKB in response to certain stimuli, released NF-KB translocates to the nucleus where it binds to specific DNA sequences and activates gene transcription. As described above, the TNFR members identified to date either include or  
40 lack an intracellular death domain region. Some TNFR molecules lacking a death domain, such as TNFR2, CD40, HVEM, and GITR, are capable of modulating NF-KB activity. [see, e.g., Lotz et al., J. Leukocyte Biol., 60:1-7 (1996)].

For a review of the TNF family of cytokines and their receptors,  
45 see Ashkenazi and Dixit, Science, 281:1305-1308 (1998); Golstein, Curr.

Biol., 7:750-753 (1997); Gruss and Dower, supra, and Nagata, Cell, 88:355-365 (1997).

#### SUMMARY OF THE INVENTION

Applicants have surprisingly found that Apo-2 ligand or other Apo-2L receptor agonists and CPT-11 can act synergistically to induce apoptosis in mammalian cells, particularly in mammalian cancer cells.

The invention provides various methods for the use of Apo-2 ligand and CPT-11 to induce apoptosis in mammalian cells. For example, the invention provides methods for inducing apoptosis comprising exposing a mammalian cell, such as a cancer cell, to CPT-11 and one or more Apo-2 ligand receptor agonists wherein CPT-11 is administered prior to the Apo-2 ligand receptor agonist(s) to pre-treat the cells.

The cells may be in cell culture or in a mammal, e.g. a mammal suffering from cancer or a condition in which induction of apoptosis in the cells is desirable. Thus, the invention includes methods for treating a mammal suffering from cancer comprising administering an effective amount of Apo-2 ligand and CPT-11, as disclosed herein.

Optionally, the methods may employ agonistic anti-Apo-2 ligand receptor antibody(s) which mimics the apoptotic activity of Apo-2 ligand. Thus, the invention provides various methods for the use of Apo-2 ligand receptor agonist antibody(s) and CPT-11 to induce apoptosis in mammalian cells. In a preferred embodiment, the agonist antibody will comprise a monoclonal antibody against the DR4 or DR5 receptor.

In optional embodiments, there are provided methods of enhancing apoptosis in mammalian cancer cells, comprising exposing mammalian cancer cells to an effective amount of CPT-11 and Apo-2 ligand receptor agonist, wherein said mammalian cancer cells are exposed to the CPT-11 about 6 hours to about 72 hours prior to exposure to said Apo-2 ligand receptor agonist. The methods may comprise exposure of said mammalian cancer cells to an effective amount of CPT-11 which induces upregulation of DR4 receptor or DR5 receptor in said cells. Optionally, the mammalian cancer cells are exposed to CPT-11 about 24 or 48 hours prior to exposure to said Apo-2 ligand receptor agonist. The Apo-2 ligand receptor agonist optionally comprises Apo2L polypeptide or anti-DR4 receptor antibody or anti-DR5 receptor antibody.

In optional embodiments, there are provided methods of treating cancer in a mammal, comprising administering to a mammal having cancer an effective amount of CPT-11 and Apo-2 ligand receptor agonist, wherein said CPT-11 is administered about 6 hours to about 72 hours prior to administration of the Apo-2 ligand receptor agonist. Optionally, the Apo-2 ligand receptor agonist comprises Apo2L polypeptide, anti-DR4 receptor antibody, or anti-DR5 receptor antibody.

The invention also provides compositions which comprise Apo-2 ligand or Apo-2L receptor agonist antibody and/or CPT-11. Optionally, the compositions of the invention will include pharmaceutically

acceptable carriers or diluents. Preferably, the compositions will include Apo-2 ligand or agonist antibody and/or CPT-11 in an amount which is effective to synergistically induce apoptosis in mammalian cells.

The invention also provides articles of manufacture and kits which  
5 include Apo-2 ligand or Apo-2L receptor agonist antibody and/or CPT-11.

#### BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows the effect of Apo-2L (open triangles), CPT-11 (open squares), Apo-2L plus CPT-11 (closed triangles), or vehicle alone (open  
10 circles) on growth of human colon carcinoma cells injected subcutaneously into athymic nude mice.

Figure 2 shows the effect of Apo-2L (60 mg/kg) (open squares), CPT-11 (80 mg/kg) (closed triangles), Apo-2L ("Apo2L.0") plus CPT-11 (closed squares), anti-DR4 mAb 4H6 (open triangles), anti-DR4 mAb plus CPT-11  
15 (closed triangles) or vehicle alone (closed circles) on growth of human colon carcinoma cells injected subcutaneously into athymic nude mice.

Figures 3A-3H show the fluorescent characterization of dead or alive tumor cells. HCT116 cultures were treated with Apo2L/TRAIL, CPT-11, and Apo2L/TRAIL + CPT-11 for 2 and 24 hours, respectively. Following  
20 incubation for 30 minutes at room temperature with the fluorescent dyes, the cells were examined with a fluorescent microscope. Calcein-positive cells (green labeling) indicate alive cells, whereas positive staining with ethidium homodimer-1 (red fluorescence) represents dead or severely damaged cells in Figures 3E, 3F, and 3G.

Figure 4 shows that CPT-11 enhanced Apo2L/TRAIL-mediated apoptosis in vitro. HCT116 cultures were incubated with CPT-11 (50 µg/ml), Apo2L/TRAIL (1 µg/ml), and Apo2L/TRAIL + CPT-11 for 24 hours. The number  
25 of live cells was determined by an alamarBlue assay (mean + SD, n = 2). The percentage of surviving cells in the sample was normalized to the control treatment.

Figures 5A-5D show how the CPT-11 sensitization of Apo2L/TRAIL induced caspase-3 activity is time dependent. HCT116 cells were treated for 2 and 24 hours with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), and Apo2L/TRAIL + CPT-11. Equivalent aliquots of cell lysates were assessed  
35 for pro-caspase-3 processing by western blot analysis (5A and 5B) and for caspase-3 activity by a fluorometric assay (5C and 5D).

Figures 6A-6B show HCT116 changes in DR5 (6A) and DR4 (6B) gene expression following treatment with Apo2L/TRAIL and CPT-11 alone or in combination. DR5 and DR4 mRNA levels were determined by bDNA assay  
40 following incubation with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), Apo2L/TRAIL + CPT-11 for 2, 6 and 24 hours, respectively. Relative values were calculated as ratios to GAPDH and normalized to untreated control cultures.

Figures 7A-7B show that Z-VAD did not block DR5 (7A) and DR4 (7B)  
45 induction in HCT116 cells following treatment with Apo2L/TRAIL and CPT-11

alone or in combination. DR5 and DR4 mRNA levels were determined by bDNA assay following incubation with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), Apo2L/TRAIL + CPT-11 for 2, 6 and 24 hours, respectively. Relative values were calculated as ratios to GAPDH and normalized to untreated control cultures.

Figure 8 shows that CPT-11 (but not Apo2L/TRAIL treatment) results in an increase in p53 protein levels in HCT116 and HUVEC cells. p53 protein levels were characterized by western blot analysis on HCT116 and HUVEC cell cultures treated for 2 and 24 hours.

Figure 9 shows that Apo2L/TRAIL treatment suppresses CPT-11-mediated induction of p21 protein levels in tumor cells but not HUVEC cells. Colon human HCT116 tumor and normal HUVEC cells were treated for 2 and 24 hours with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), and Apo2L/TRAIL + CPT-11. Equivalent aliquots of cell lysates (50 µg/lane) were tested for p21 protein expression by western blot analysis.

Figure 10 shows that the caspase-8 inhibitor FLIP protein levels did not change after treatments. Colon human HCT116 tumor cells were treated for 2 and 24 hours with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), and Apo2L/TRAIL + CPT-11 (1 µg/ml). Following treatment, the cell cultures were processed by flow cytometric cell cycle analysis.

Figure 11 shows that Apo2L/TRAIL suppresses CPT-11 induced G2/M cell cycle arrest. HCT116 tumor cells were treated for 2, 6 and 24 hours with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), and Apo2L/TRAIL + CPT-11. Following treatment, the cell cultures were processed by flow cytometric cell cycle analysis.

Figure 12 shows that the caspase inhibitor Z-VAD (I) has a differential effect on the levels of p53 and p21. Cell cultures were treated with or without 20 µM of Z-VAD for 24 hours. Cell lysates were processed for western blot analysis of p53 and p21 protein levels.

Figure 13 shows that the Apo2L/TRAIL treatment in the presence of the caspase inhibitor ZVAD fails to prevent the CPT-11 induced G2-M arrest. HCT116 tumor cells were treated for 24 hours with Apo2L/TRAIL (1 µg/ml), CPT-11 (50 µg/ml), and Apo2L/TRAIL + CPT-11 in the presence of 20 µM of Z-VAD. Following treatment, the cell cultures were processed by flow cytometric cell cycle analysis.

Figures 14A-14B show that sequential treatment enhances total cell killing in tumor cells. A. In the combination group, HCT116 cells were exposed to CPT-11 (10 microgram/ml) and different concentrations of Apo2L/TRAIL (as indicated in the figure) for a total of 24 hours, followed by another 24 hours of incubation in the presence of medium alone. In the sequential group, cells were exposed for the initial 24 hours to CPT-11, then changed to Apo2L/TRAIL containing medium for another 24 hours. Cell survival was determined by the crystal violet

assay as described in the Examples (mean  $\pm$  SD, n = 4). B. HCT116 cells (combination) were exposed to CPT-11 (10 microgram/ml) and different concentrations of Apo2L/TRAIL for a total of 24 hours, followed by another 120 hours of incubation in the presence of medium alone. In the sequential group, cells were exposed for the initial 24 hours to CPT-11, then changed to Apo2L/TRAIL containing medium for another 24 hours followed by incubation of drug free medium for 96 hours and tested for cell survival as before.

Figure 15 shows that substitution of SN-38 instead of CPT-11 in the combination and sequential treatment results in similar enhanced tumor cell killing. In the combination group, HCT116 cells were exposed to SN-38 (0.05 microgram/ml) and two concentrations of Apo2L/TRAIL (as indicated in the figure) for a total of 24 hours, followed by another 24 hours of incubation in the presence of medium alone. In the sequential group, cells were exposed for the initial 24 hours to SN-38, then changed to Apo2L/TRAIL containing medium alone for another 24 hours. Cell survival was determined by crystal violet assay as described in the Examples (mean  $\pm$  SD, n = 4).

20

#### DETAILED DESCRIPTION OF THE INVENTION

##### **I. Definitions**

The terms "apoptosis" and "apoptotic activity" are used in a broad sense and refer to the orderly or controlled form of cell death in mammals that is typically accompanied by one or more characteristic cell changes, including condensation of cytoplasm, loss of plasma membrane microvilli, segmentation of the nucleus, degradation of chromosomal DNA or loss of mitochondrial function. This activity can be determined and measured using techniques known in the art, for instance, by cell viability assays, FACS analysis or DNA electrophoresis, and more specifically by binding of annexin V, fragmentation of DNA, PARP cleavage, cell shrinkage, dilation of endoplasmic reticulum, cell fragmentation, and/or formation of membrane vesicles (called apoptotic bodies). These techniques and assays are described in the art, for example, in WO97/25428 and WO97/01633. Optionally, apoptotic activity may be measured using the assays described in the Examples.

As used herein, the term "synergy" or "synergism" or "synergistically" refers to the interaction of two or more agents so that their combined effect is greater than the sum of their individual effects.

The terms "Apo-2 ligand", "Apo-2L", or "TRAIL" are used herein to refer to a polypeptide which includes amino acid residues 95-281, inclusive, 114-281, inclusive, residues 91-281, inclusive, residues 92-281, inclusive, residues 41-281, inclusive, residues 15-281, inclusive, or residues 1-281, inclusive, of the amino acid sequence shown in Figure 1A

of Pitti et al., J. Biol. Chem., 271:12687-12690 (1996) (provided herein in the Sequence Listing as SEQ ID NO:1), as well as biologically active (e.g., having apoptotic activity) fragments, deletional, insertional, or substitutional variants of the above sequences. In one embodiment, the polypeptide sequence comprises residues 114-281 of SEQ ID NO:1. Optionally, the polypeptide sequence has at least residues 91-281 or residues 92-281 of SEQ ID NO:1. In another preferred embodiment, the biologically active fragments or variants have at least about 80% amino acid sequence identity, more preferably at least about 90% amino acid sequence identity, and even more preferably, at least about 95%, 96%, 97%, 98%, or 99% amino acid sequence identity with any one of the above sequences. The definition encompasses substitutional variants of the Apo-2 ligand comprising amino acids 91-281 of Figure 1A of Pitti et al., J. Biol. Chem., 271:12687-12690 (1996) (SEQ ID NO:1) in which at least one of the amino acids at positions 203, 218 or 269 (using the numbering of the sequence provided in Pitti et al., supra (SEQ ID NO:1)) are substituted by an alanine residue. The definition encompasses Apo-2 ligand isolated from an Apo-2 ligand source, such as from human tissue types, or from another source, or prepared by recombinant or synthetic methods. The term Apo-2 ligand also refers to the polypeptides described in WO 97/25428, supra, and WO97/01633, supra. It is contemplated that the Apo-2 ligand polypeptide may be linked to one or more polymer molecules such as polyethylene glycol.

The term "CPT-11" is used in a general sense and refers to a chemotherapy or chemotherapeutic agent which is of the topoisomerase I inhibitor class. The term "CPT-11" as used herein includes the chemotherapeutic agents having the chemical name (4S)-4, 11-diethyl-4-hydroxy-9-[(4-piperidino-piperidino)carbonyloxyl]-1H-pyrano[3', 4':6,7]indolizino [1, 2-b]quinoline-3, 14(4H, 12H)dione hydrochloride trihydrate, and the names irinotecan, camptothecin, topotecan, or Camptosar®, as well as water-soluble derivatives thereof or pharmaceutically acceptable salts of such agents. Irinotecan hydrochloride has the empirical formula  $C_{33}H_{38}N_4O_6 \cdot HCl \cdot 3H_2O$  and a molecular weight of approximately 677.19. Such chemical names and chemical formulae will be readily familiar to those skilled in the art. Camptosar® is commercially available from Pharmacia & Upjohn and approved for marketing in the United States by the FDA. The product insert for Camptosar® indicates the molecule can be used for treatment of human patients with metastatic colorectal carcinoma whose disease has recurred or progressed following 5-FU based therapy. It is contemplated that the CPT-11 may be linked to one or more polymer molecules such as polyethylene glycol.

"Percent (%) amino acid sequence identity" with respect to the Apo-2L polypeptide sequences identified herein is defined as the percentage of amino acid residues in a candidate sequence that are identical with the amino acid residues in an Apo-2L sequence, after aligning the sequences



and introducing gaps, if necessary, to achieve the maximum percent sequence identity, and not considering any conservative substitutions as part of the sequence identity. Alignment for purposes of determining percent amino acid sequence identity can be achieved in various ways that are within the skill in the art, for instance, using publicly available computer software such as BLAST, BLAST-2, ALIGN, ALIGN-2 or Megalign (DNASTAR) software. Those skilled in the art can determine appropriate parameters for measuring alignment, including any algorithms needed to achieve maximal alignment over the full-length of the sequences being compared. Optionally, % amino acid sequence identity values are obtained by using the sequence comparison computer program ALIGN-2. The ALIGN-2 sequence comparison computer program was authored by Genentech, Inc. and the source code has been filed with user documentation in the U.S. Copyright Office, Washington, D.C., 20559, where it is registered under U.S. Copyright Registration No. TXU510087. The ALIGN-2 program is publicly available through Genentech, Inc., South San Francisco, California. The ALIGN-2 program should be compiled for use on a UNIX operating system, preferably digital UNIX V4.0D. All sequence comparison parameters are set by the ALIGN-2 program and do not vary. However, % amino acid sequence identity may also be determined using the sequence comparison program NCBI-BLAST2 (Altschul et al., Nucleic Acids Res. 25:3389-3402 (1997)). The NCBI-BLAST2 sequence comparison program may be downloaded from <http://www.ncbi.nlm.nih.gov>. NCBI-BLAST2 uses several search parameters, wherein all of those search parameters are set to default values including, for example, unmask = yes, strand = all, expected occurrences = 10, minimum low complexity length = 15/5, multi-pass e-value = 0.01, constant for multi-pass = 25, dropoff for final gapped alignment = 25 and scoring matrix = BLOSUM62.

The term "antibody" when used in reference to an "agonistic anti-Apo-2 ligand receptor antibody" is used in the broadest sense and specifically covers intact monoclonal antibodies, polyclonal antibodies, multispecific antibodies (e.g. bispecific antibodies) formed from at least two intact antibodies, and antibody fragments so long as they bind one or more Apo-2 ligand receptors and/or are capable of activating the apoptosis signaling pathway of the mammalian cell expressing one or more of the Apo-2 ligand receptors or mimic (e.g., have comparable or at least equal to) the apoptotic activity of Apo-2 ligand or have greater apoptotic activity than that of Apo-2 ligand.

"Apo-2 ligand receptor" includes the receptors referred to in the art as "DR4" and "DR5". Pan et al. have described the TNF receptor family member referred to as "DR4" [Pan et al., Science, 276:111-113 (1997); see also WO98/32856 published July 30, 1998]. The DR4 receptor was reported to contain a cytoplasmic death domain capable of engaging the cell suicide apparatus. Pan et al. disclose that DR4 is believed to be a receptor for the ligand known as Apo2L/TRAIL. The amino acid

sequence of the full length DR4 receptor is provided herein in SEQ ID NO:2. Sheridan et al., Science, 277:818-821 (1997) and Pan et al., Science, 277:815-818 (1997) described another receptor for Apo2L/TRAIL [see also, WO98/51793 published November 19, 1998; WO98/41629 published September 24, 1998]. This receptor is referred to as DR5 (the receptor has also been alternatively referred to as Apo-2; TRAIL-R, TR6, Tango-63, hAPO8, TRICK2 or KILLER; Screaton et al., Curr. Biol., 7:693-696 (1997); Walczak et al., EMBO J., 16:5386-5387 (1997); Wu et al., Nature Genetics, 17:141-143 (1997); WO98/35986 published August 20, 1998 (corresponding to issued US Patent 6,072,047); EP870,827 published October 14, 1998; WO98/46643 published October 22, 1998; WO99/02653 published January 21, 1999; WO99/09165 published February 25, 1999; WO99/11791 published March 11, 1999]. Like DR4, DR5 is reported to contain a cytoplasmic death domain and be capable of signaling apoptosis. The full length DR5 receptor sequence in WO98/35986 (corresponding to US Patent 6,072,047) is reported to be a 440 amino acid polypeptide, and that amino acid sequence is provided in SEQ ID NO:3. The full length DR5 receptor sequence in WO98/51793 is reported to be a 411 amino acid polypeptide, and that amino acid sequence is provided in SEQ ID NO:4. As described above, other receptors for Apo-2L include DcR1, DcR2, and OPG [see, Sheridan et al., supra; Marsters et al., supra; and Simonet et al., supra]. The term "Apo-2L receptor" when used herein encompasses native sequence receptor and receptor variants. These terms encompass Apo-2L receptor expressed in a variety of mammals, including humans. Apo-2L receptor may be endogenously expressed as occurs naturally in a variety of human tissue lineages, or may be expressed by recombinant or synthetic methods. A "native sequence Apo-2L receptor" comprises a polypeptide having the same amino acid sequence as an Apo-2L receptor derived from nature. Thus, a native sequence Apo-2L receptor can have the amino acid sequence of naturally-occurring Apo-2L receptor from any mammal. Such native sequence Apo-2L receptor can be isolated from nature or can be produced by recombinant or synthetic means. The term "native sequence Apo-2L receptor" specifically encompasses naturally-occurring truncated or secreted forms of the receptor (e.g., a soluble form containing, for instance, an extracellular domain sequence), naturally-occurring variant forms (e.g., alternatively spliced forms) and naturally-occurring allelic variants. Receptor variants may include fragments or deletion mutants of the native sequence Apo-2L receptor.

The term "monoclonal antibody" as used herein refers to an antibody obtained from a population of substantially homogeneous antibodies, i.e., the individual antibodies comprising the population are identical except for possible naturally occurring mutations that may be present in minor amounts. Monoclonal antibodies are highly specific, being directed against a single antigenic site. Furthermore, in contrast to conventional (polyclonal) antibody preparations which typically include

different antibodies directed against different determinants (epitopes), each monoclonal antibody is directed against a single determinant on the antigen. In addition to their specificity, the monoclonal antibodies are advantageous in that they are synthesized by the hybridoma culture, uncontaminated by other immunoglobulins. The modifier "monoclonal" indicates the character of the antibody as being obtained from a substantially homogeneous population of antibodies, and is not to be construed as requiring production of the antibody by any particular method. For example, the monoclonal antibodies to be used in accordance with the present invention may be made by the hybridoma method first described by Kohler et al., Nature, 256:495 (1975), or may be made by recombinant DNA methods (see, e.g., U.S. Patent No. 4,816,567). The "monoclonal antibodies" may also be isolated from phage antibody libraries using the techniques described in Clackson et al., Nature, 352:624-628 (1991) and Marks et al., J. Mol. Biol., 222:581-597 (1991), for example.

The monoclonal antibodies herein specifically include "chimeric" antibodies (immunoglobulins) in which a portion of the heavy and/or light chain is identical with or homologous to corresponding sequences in antibodies derived from a particular species or belonging to a particular antibody class or subclass, while the remainder of the chain(s) is identical with or homologous to corresponding sequences in antibodies derived from another species or belonging to another antibody class or subclass, as well as fragments of such antibodies, so long as they exhibit the desired biological activity (U.S. Patent No. 4,816,567; Morrison et al., Proc. Natl. Acad. Sci. USA, 81:6851-6855 (1984)).

"Humanized" forms of non-human (e.g., murine) antibodies are chimeric immunoglobulins, immunoglobulin chains or fragments thereof (such as Fv, Fab, Fab', F(ab')<sub>2</sub> or other antigen-binding subsequences of antibodies) which contain minimal sequence derived from non-human immunoglobulin. For the most part, humanized antibodies are human immunoglobulins (recipient antibody) in which residues from a complementarity-determining region (CDR) of the recipient are replaced by residues from a CDR of a non-human species (donor antibody) such as mouse, rat or rabbit having the desired specificity, affinity, and capacity. In some instances, Fv framework region (FR) residues of the human immunoglobulin are replaced by corresponding non-human residues. Furthermore, humanized antibodies may comprise residues which are found neither in the recipient antibody nor in the imported CDR or framework sequences. These modifications are made to further refine and maximize antibody performance. In general, the humanized antibody will comprise substantially all of at least one, and typically two, variable domains, in which all or substantially all of the CDR regions correspond to those of a non-human immunoglobulin and all or substantially all of the FR regions are those of a human immunoglobulin sequence. The humanized

antibody optimally also will comprise at least a portion of an immunoglobulin constant region (Fc), typically that of a human immunoglobulin. For further details, see Jones et al., Nature, 321:522-525 (1986); Reichmann et al., Nature, 332:323-329 (1988); and Presta, 5 Curr. Op. Struct. Biol., 2:593-596 (1992). The humanized antibody includes a PRIMATIZED™ antibody wherein the antigen-binding region of the antibody is derived from an antibody produced by immunizing macaque monkeys with the antigen of interest.

Antibodies are typically proteins or polypeptides which exhibit 10 binding specificity to a specific antigen. Native antibodies are usually heterotetrameric glycoproteins, composed of two identical light (L) chains and two identical heavy (H) chains. Typically, each light chain is linked to a heavy chain by one covalent disulfide bond, while the number of disulfide linkages varies between the heavy chains of different 15 immunoglobulin isotypes. Each heavy and light chain also has regularly spaced intrachain disulfide bridges. Each heavy chain has at one end a variable domain (V<sub>H</sub>) followed by a number of constant domains. Each light chain has a variable domain at one end (V<sub>L</sub>) and a constant domain at its other end; the constant domain of the light chain is aligned with the 20 first constant domain of the heavy chain, and the light chain variable domain is aligned with the variable domain of the heavy chain. Particular amino acid residues are believed to form an interface between the light and heavy chain variable domains [Chothia et al., J. Mol. Biol., 186:651-663 (1985); Novotny and Haber, Proc. Natl. Acad. Sci. USA, 82:4592-4596 25 (1985)]. The light chains of antibodies from any vertebrate species can be assigned to one of two clearly distinct types, called kappa and lambda, based on the amino acid sequences of their constant domains. Depending on the amino acid sequence of the constant domain of their heavy chains, immunoglobulins can be assigned to different classes. There are five 30 major classes of immunoglobulins: IgA, IgD, IgE, IgG and IgM, and several of these may be further divided into subclasses (isotypes), e.g., IgG-1, IgG-2, IgG-3, and IgG-4; IgA-1 and IgA-2. The heavy chain constant domains that correspond to the different classes of immunoglobulins are called alpha, delta, epsilon, gamma, and mu, respectively.

35 "Antibody fragments" comprise a portion of an intact antibody, generally the antigen binding or variable region of the intact antibody. Examples of antibody fragments include Fab, Fab', F(ab')<sub>2</sub>, and Fv fragments, diabodies, single chain antibody molecules, and multispecific antibodies formed from antibody fragments.

40 The term "variable" is used herein to describe certain portions of the variable domains which differ in sequence among antibodies and are used in the binding and specificity of each particular antibody for its particular antigen. However, the variability is not usually evenly distributed through the variable domains of antibodies. It is typically 45 concentrated in three segments called complementarity determining regions

(CDRs) or hypervariable regions both in the light chain and the heavy chain variable domains. The more highly conserved portions of the variable domains are called the framework (FR). The variable domains of native heavy and light chains each comprise four FR regions, largely adopting a  $\beta$ -sheet configuration, connected by three CDRs, which form loops connecting, and in some cases forming part of, the  $\beta$ -sheet structure. The CDRs in each chain are held together in close proximity by the FR regions and, with the CDRs from the other chain, contribute to the formation of the antigen binding site of antibodies [see Kabat, E.A. et al., Sequences of Proteins of Immunological Interest, National Institutes of Health, Bethesda, MD (1987)]. The constant domains are not involved directly in binding an antibody to an antigen, but exhibit various effector functions, such as participation of the antibody in antibody-dependent cellular toxicity.

The monoclonal antibodies herein include chimeric, hybrid and recombinant antibodies produced by splicing a variable (including hypervariable) domain of an anti-Apo-2L receptor antibody with a constant domain (e.g. "humanized" antibodies), or a light chain with a heavy chain, or a chain from one species with a chain from another species, or fusions with heterologous proteins, regardless of species of origin or immunoglobulin class or subclass designation, as well as antibody fragments (e.g., Fab, F(ab')<sub>2</sub>, and Fv), so long as they exhibit the desired biological activity or properties. See, e.g. U.S. Pat. No. 4,816,567 and Mage et al., in Monoclonal Antibody Production Techniques and Applications, pp.79-97 (Marcel Dekker, Inc.: New York, 1987).

A "human antibody" is one which possesses an amino acid sequence which corresponds to that of an antibody produced by a human and/or has been made using any of the techniques for making human antibodies as disclosed herein. This definition of a human antibody specifically excludes a humanized antibody comprising non-human antigen-binding residues. Human antibodies can be produced using various techniques known in the art. In one embodiment, the human antibody is selected from a phage library, where that phage library expresses human antibodies (Vaughan et al. Nature Biotechnology, 14:309-314 (1996); Sheets et al. PNAS, (USA) 95:6157-6162 (1998)); Hoogenboom and Winter, J. Mol. Biol., 227:381 (1991); Marks et al., J. Mol. Biol., 222:581 (1991)). Human antibodies can also be made by introducing human immunoglobulin loci into transgenic animals, e.g., mice in which the endogenous immunoglobulin genes have been partially or completely inactivated. Upon challenge, human antibody production is observed, which closely resembles that seen in humans in all respects, including gene rearrangement, assembly, and antibody repertoire. This approach is described, for example, in U.S. Patent Nos. 5,545,807; 5,545,806; 5,569,825; 5,625,126; 5,633,425; 5,661,016, and in the following scientific publications: Marks et al.,

Bio/Technology, 10: 779-783 (1992); Lonberg et al., Nature, 368: 856-859 (1994); Morrison, Nature, 368:812-13 (1994); Fishwild et al., Nature Biotechnology, 14: 845-51 (1996); Neuberger, Nature Biotechnology, 14: 826 (1996); Lonberg and Huszar, Intern. Rev. Immunol., 13:65-93 (1995).  
5 Alternatively, the human antibody may be prepared via immortalization of human B lymphocytes producing an antibody directed against a target antigen (such B lymphocytes may be recovered from an individual or may have been immunized *in vitro*). See, e.g., Cole et al., Monoclonal Antibodies and Cancer Therapy, Alan R. Liss, p. 77 (1985); Boerner et al., J. Immunol., 147 (1):86-95 (1991); and US Pat No. 5,750,373.

The term "Fc region" is used to define the C-terminal region of an immunoglobulin heavy chain which may be generated by papain digestion of an intact antibody. The Fc region may be a native sequence Fc region or a variant Fc region. Although the boundaries of the Fc region of an immunoglobulin heavy chain might vary, the human IgG heavy chain Fc region is usually defined to stretch from an amino acid residue at about position Cys226, or from about position Pro230, to the carboxyl-terminus of the Fc region (using herein the numbering system according to Kabat et al., supra). The Fc region of an immunoglobulin generally comprises two constant domains, a CH2 domain and a CH3 domain, and optionally comprises a CH4 domain.  
15

By "Fc region chain" herein is meant one of the two polypeptide chains of an Fc region.

The "CH2 domain" of a human IgG Fc region (also referred to as "C $\gamma$ 2" domain) usually extends from an amino acid residue at about position 231 to an amino acid residue at about position 340. The CH2 domain is unique in that it is not closely paired with another domain. Rather, two N-linked branched carbohydrate chains are interposed between the two CH2 domains of an intact native IgG molecule. It has been speculated that the carbohydrate may provide a substitute for the domain-domain pairing and help stabilize the CH2 domain. Burton, Molec. Immunol. 22:161-206 (1985). The CH2 domain herein may be a native sequence CH2 domain or variant CH2 domain.  
25  
30

The "CH3 domain" comprises the stretch of residues C-terminal to a CH2 domain in an Fc region (i.e. from an amino acid residue at about position 341 to an amino acid residue at about position 447 of an IgG). The CH3 region herein may be a native sequence CH3 domain or a variant CH3 domain (e.g. a CH3 domain with an introduced "protuberance" in one chain thereof and a corresponding introduced "cavity" in the other chain thereof; see US Patent No. 5,821,333).  
35  
40

"Hinge region" is generally defined as stretching from about Glu216, or about Cys226, to about Pro230 of human IgG1 (Burton, Molec. Immunol. 22:161-206 (1985)). Hinge regions of other IgG isotypes may be aligned with the IgG1 sequence by placing the first and last cysteine residues forming inter-heavy chain S-S bonds in the same positions. The  
45

hinge region herein may be a native sequence hinge region or a variant hinge region. The two polypeptide chains of a variant hinge region generally retain at least one cysteine residue per polypeptide chain, so that the two polypeptide chains of the variant hinge region can form a disulfide bond between the two chains. The preferred hinge region herein is a native sequence human hinge region, e.g. a native sequence human IgG1 hinge region.

A "functional Fc region" possesses at least one "effector function" of a native sequence Fc region. Exemplary "effector functions" include Clq binding; complement dependent cytotoxicity (CDC); Fc receptor binding; antibody-dependent cell-mediated cytotoxicity (ADCC); phagocytosis; down regulation of cell surface receptors (e.g. B cell receptor; BCR), etc. Such effector functions generally require the Fc region to be combined with a binding domain (e.g. an antibody variable domain) and can be assessed using various assays known in the art for evaluating such antibody effector functions.

A "native sequence Fc region" comprises an amino acid sequence identical to the amino acid sequence of an Fc region found in nature. A "variant Fc region" comprises an amino acid sequence which differs from that of a native sequence Fc region by virtue of at least one amino acid modification. Preferably, the variant Fc region has at least one amino acid substitution compared to a native sequence Fc region or to the Fc region of a parent polypeptide, e.g. from about one to about ten amino acid substitutions, and preferably from about one to about five amino acid substitutions in a native sequence Fc region or in the Fc region of the parent polypeptide. The variant Fc region herein will preferably possess at least about 80% sequence identity with a native sequence Fc region and/or with an Fc region of a parent polypeptide, and most preferably at least about 90% sequence identity therewith, more preferably at least about 95% sequence identity therewith.

The terms "Fc receptor" and "FcR" are used to describe a receptor that binds to the Fc region of an antibody. The preferred FcR is a native sequence human FcR. Moreover, a preferred FcR is one which binds an IgG antibody (a gamma receptor) and includes receptors of the FcγRI, FcγRII, and FcγRIII subclasses, including allelic variants and alternatively spliced forms of these receptors. FcγRII receptors include FcγRIIA (an "activating receptor") and FcγRIIB (an "inhibiting receptor"), which have similar amino acid sequences that differ primarily in the cytoplasmic domains thereof. Activating receptor FcγRIIA contains an immunoreceptor tyrosine-based activation motif (ITAM) in its cytoplasmic domain. Inhibiting receptor FcγRIIB contains an immunoreceptor tyrosine-based inhibition motif (ITIM) in its cytoplasmic domain (reviewed in Daëron, Annu. Rev. Immunol., 15:203-234 (1997)). FcRs are reviewed in Ravetch and Kinet, Annu. Rev. Immunol., 9:457-92

(1991); Capel et al., Immunomethods, 4:25-34 (1994); and de Haas et al., J. Lab. Clin. Med., 126:330-41 (1995). Other FcRs, including those to be identified in the future, are encompassed by the term "FcR" herein. The term also includes the neonatal receptor, FcRn, which is responsible for the transfer of maternal IgGs to the fetus (Guyer et al., J. Immunol., 117:587 (1976); and Kim et al., J. Immunol., 24:249 (1994)).

An "affinity matured" antibody is one with one or more alterations in one or more CDRs thereof which result in an improvement in the affinity of the antibody for antigen, compared to a parent antibody which does not possess those alteration(s). Preferred affinity matured antibodies will have nanomolar or even picomolar affinities for the target antigen. Affinity matured antibodies are produced by procedures known in the art. Marks et al. Bio/Technology, 10:779-783 (1992) describes affinity maturation by VH and VL domain shuffling. Random mutagenesis of CDR and/or framework residues is described by: Barbas et al. Proc Nat. Acad. Sci, USA 91:3809-3813 (1994); Schier et al. Gene, 169:147-155 (1995); Yelton et al. J. Immunol., 155:1994-2004 (1995); Jackson et al., J. Immunol., 154(7):3310-9 (1995); and Hawkins et al, J. Mol. Biol., 226:889-896 (1992).

The terms "agonist" and "agonistic" when used herein refer to or describe a molecule which is capable of, directly or indirectly, substantially inducing, promoting or enhancing biological activity or activation of a receptor for Apo-2 ligand. Optionally, an "agonist Apo-2L receptor antibody" is an antibody which has activity that mimics or is comparable to Apo-2 ligand. Preferably, the agonist is a molecule which is capable of inducing apoptosis in a mammalian cell, preferably, a mammalian cancer cell. Even more preferably, the agonist is an antibody directed to an Apo-2L receptor and said antibody has apoptotic activity which is equal to or greater than the Apo-2L polypeptide described in Example 1. Optionally, the agonist activity of such molecule can be determined by assaying the molecule, alone or in a cross-linked form using Fc immunoglobulin or complement (described below), in an assay described in Example 2 to examine apoptosis of 9D cells or other cells which express a receptor for Apo-2L such as DR4 or DR5. It is contemplated that the agonist may be linked to one or more polymer molecules such as polyethylene glycol.

"Isolated," when used to describe the various proteins disclosed herein, means protein that has been identified and separated and/or recovered from a component of its natural environment. Contaminant components of its natural environment are materials that would typically interfere with diagnostic or therapeutic uses for the protein, and may include enzymes, hormones, and other proteinaceous or non-proteinaceous solutes. In preferred embodiments, the protein will be purified (1) to a degree sufficient to obtain at least 15 residues of N-terminal or internal amino acid sequence by use of a spinning cup sequenator, or (2) to



homogeneity by SDS-PAGE under non-reducing or reducing conditions using Coomassie blue or, preferably, silver stain. Isolated protein includes protein *in situ* within recombinant cells, since at least one component of the protein natural environment will not be present. Ordinarily, however, isolated protein will be prepared by at least one purification step.

"Biologically active" or "biological activity" for the purposes herein means (a) having the ability to induce or stimulate apoptosis in at least one type of mammalian cell (such as a cancer cell) or virally-infected cell *in vivo* or *ex vivo*; (b) capable of raising an antibody, i.e., immunogenic; or (c) retaining the activity of a native or naturally-occurring Apo-2 ligand polypeptide.

A "growth inhibitory agent" when used herein refers to a compound or composition which inhibits growth of a cell *in vitro* and/or *in vivo*. Thus, the growth inhibitory agent may be one which significantly reduces the percentage of cells in S phase. Examples of growth inhibitory agents include agents that block cell cycle progression (at a place other than S phase), such as agents that induce G1 arrest and M-phase arrest. Classical M-phase blockers include the vincas (vincristine and vinblastine), TAXOL®, and topo II inhibitors such as doxorubicin, epirubicin, daunorubicin, etoposide, and bleomycin. Those agents that arrest G1 also spill over into S-phase arrest, for example, DNA alkylating agents such as tamoxifen, prednisone, dacarbazine, mechlorethamine, cisplatin, methotrexate, 5-fluorouracil, and ara-C. Further information can be found in The Molecular Basis of Cancer, Mendelsohn and Israel, eds., Chapter 1, entitled "Cell cycle regulation, oncogenes, and antineoplastic drugs" by Murakami et al. (WB Saunders: Philadelphia, 1995), especially p. 13.

The term "prodrug" as used in this application refers to a precursor or derivative form of a pharmaceutically active substance that is less cytotoxic to cancer cells compared to the parent drug and is capable of being enzymatically activated or converted into the more active parent form. See, e.g., Wilman, "Prodrugs in Cancer Chemotherapy" *Biochemical Society Transactions*, 14, pp. 375-382, 615th Meeting Belfast (1986) and Stella et al., "Prodrugs: A Chemical Approach to Targeted Drug Delivery," *Directed Drug Delivery*, Borchardt et al., (ed.), pp. 247-267, Humana Press (1985). The prodrugs of this invention include, but are not limited to, phosphate-containing prodrugs, thiophosphate-containing prodrugs, sulfate-containing prodrugs, peptide-containing prodrugs, D-amino acid-modified prodrugs, glycosylated prodrugs, beta-lactam-containing prodrugs, optionally substituted phenoxyacetamide-containing prodrugs or optionally substituted phenylacetamide-containing prodrugs, 5-fluorocytosine and other 5-fluorouridine prodrugs which can be converted into the more active cytotoxic free drug. Examples of cytotoxic drugs that can be derivatized into a prodrug form for use in

this invention include, but are not limited to, those chemotherapeutic agents described below.

The term "cytotoxic agent" as used herein refers to a substance that inhibits or prevents the function of cells and/or causes destruction of cells. The term is intended to include radioactive isotopes (e.g. At<sup>211</sup>, I<sup>131</sup>, I<sup>125</sup>, Y<sup>90</sup>, Re<sup>186</sup>, Re<sup>188</sup>, Sm<sup>153</sup>, Bi<sup>212</sup>, P<sup>32</sup> and radioactive isotopes of Lu), chemotherapeutic agents, and toxins such as small molecule toxins or enzymatically active toxins of bacterial, fungal, plant or animal origin, including fragments and/or variants thereof.

A "chemotherapeutic agent" is a chemical compound useful in the treatment of conditions like cancer. Examples of chemotherapeutic agents include alkylating agents such as thiotepa and cyclophosphamide (CYTOXAN<sup>TM</sup>); alkyl sulfonates such as busulfan, improsulfan and piposulfan; aziridines such as benzodopa, carboquone, meturedopa, and uredopa; ethylenimines and methylamelamines including altretamine, triethylenemelamine, triethylenephosphoramide, triethylenethiophosphoramide and trimethylolomelamine; acetogenins (especially bullatacin and bullatacinone); a camptothecin (including the synthetic analogue topotecan); bryostatin; callystatin; CC-1065 (including its adozelesin, carzelesin and bizelesin synthetic analogues); cryptophycins (particularly cryptophycin 1 and cryptophycin 8); dolastatin; duocarmycin (including the synthetic analogues, KW-2189 and CBI-TMI); eleutherobin; pancratistatin; a sarcodictyin; spongistatin; nitrogen mustards such as chlorambucil, chlornaphazine, cholophosphamide, estramustine, ifosfamide, mechlorethamine, mechlorethamine oxide hydrochloride, melphalan, novembichin, phenesterine, prednimustine, trofosfamide, uracil mustard; nitrosureas such as carmustine, chlorozotocin, fotemustine, lomustine, nimustine, ranimustine; antibiotics such as the enediyne antibiotics (e.g. calicheamicin, especially calicheamicin  $\gamma_1^I$  and calicheamicin  $\theta_1^I$ , see, e.g., Agnew Chem Intl. Ed. Engl., 33:183-186 (1994); dynemicin, including dynemicin A; an esperamicin; as well as neocarzinostatin chromophore and related chromoprotein enediyne antibiotic chromomorphores), aclacinomysins, actinomycin, authramycin, azaserine, bleomycins, cactinomycin, carabacin, carminomycin, carzinophilin, chromomycins, dactinomycin, daunorubicin, detorubicin, 6-diazo-5-oxo-L-norleucine, doxorubicin (including morpholino-doxorubicin, cyanomorpholino-doxorubicin, 2-pyrrolino-doxorubicin and deoxydoxorubicin), epirubicin, esorubicin, idarubicin, marcellomycin, mitomycins, mycophenolic acid, nogalamycin, olivomycins, peplomycin, potfiromycin, puromycin, quelamycin, rodorubicin, streptonigrin, streptozocin, tubercidin, ubenimex, zinostatin, zorubicin; anti-metabolites such as methotrexate and 5-fluorouracil (5-FU); folic acid analogues such as denopterin, methotrexate, pteropterin, trimetrexate; purine analogs such as fludarabine, 6-mercaptopurine, thiamiprine, thioguanine; pyrimidine analogs such as ancitabine, azacitidine, 6-

azauridine, carmofur, cytarabine, dideoxyuridine, doxifluridine, enocitabine, floxuridine, 5-FU; androgens such as calusterone, dromostanolone propionate, epitio stanol, mepitio stanane, testolactone; anti-adrenals such as aminogluthethimide, mitotane, trilostane; folic acid  
5 replenisher such as frolinic acid; aceglatone; aldophosphamide glycoside; aminolevulinic acid; amsacrine; bestrabucil; bisantrene; edatraxate; defofamine; demecolcine; diaziquone; elfornithine; elliptinium acetate; an epothilone; etoglucid; gallium nitrate; hydroxyurea; lentinan; lonidamine; maytansinoids such as maytansine and ansamitocins;  
10 mitoguazone; mitoxantrone; mopidamol; nitracrine; pentostatin; phenamet; pirarubicin; podophyllinic acid; 2-ethylhydrazide; procarbazine; PSK®; razoxane; rhizoxin; sizofiran; spirogermanium; tenuazonic acid; triaziquone; 2, 2', 2''-trichlorotriethylamine; trichothecenes (especially T-2 toxin, verracurin A, roridin A and anguidine); urethan; vindesine;  
15 dacarbazine; mannomustine; mitobronitol; mitolactol; pipobroman; gacytosine; arabinoside ("Ara-C"); cyclophosphamide; thiotepea; taxoids, e.g. paclitaxel (TAXOL®, Bristol-Myers Squibb Oncology, Princeton, NJ) and doxetaxel (TAXOTERE®, Rhône-Poulenc Rorer, Antony, France); chlorambucil; gemcitabine; 6-thioguanine; mercaptopurine; methotrexate;  
20 platinum analogs such as cisplatin and carboplatin; vinblastine; platinum; etoposide (VP-16); ifosfamide; mitomycin C; mitoxantrone; vincristine; vinorelbine; navelbine; novantrone; teniposide; daunomycin; aminopterin; xeloda; ibandronate; CPT-11; topoisomerase inhibitor RFS 2000; difluoromethylornithine (DMFO); retinoic acid; capecitabine; and  
25 pharmaceutically acceptable salts, acids or derivatives of any of the above. Also included in this definition are anti-hormonal agents that act to regulate or inhibit hormone action on tumors such as anti-estrogens including for example tamoxifen, raloxifene, aromatase inhibiting 4(5)-imidazoles, 4-hydroxytamoxifen, trioxifene, keoxifene,  
30 LY117018, onapristone, and toremifene (Fareston); and anti-androgens such as flutamide, nilutamide, bicalutamide, leuprolide, and goserelin; and pharmaceutically acceptable salts, acids or derivatives of any of the above.

The term "cytokine" is a generic term for proteins released by one  
35 cell population which act on another cell as intercellular mediators. Examples of such cytokines are lymphokines, monokines, and traditional polypeptide hormones. Included among the cytokines are growth hormone such as human growth hormone, N-methionyl human growth hormone, and bovine growth hormone; parathyroid hormone; thyroxine; insulin;  
40 proinsulin; relaxin; prorelaxin; glycoprotein hormones such as follicle stimulating hormone (FSH), thyroid stimulating hormone (TSH), and luteinizing hormone (LH); hepatic growth factor; fibroblast growth factor; prolactin; placental lactogen; tumor necrosis factor-alpha and -beta; mullerian-inhibiting substance; mouse gonadotropin-associated  
45 peptide; inhibin; activin; vascular endothelial growth factor; integrin;

thrombopoietin (TPO); nerve growth factors such as NGF-alpha; platelet-growth factor; transforming growth factors (TGFs) such as TGF-alpha and TGF-beta; insulin-like growth factor-I and -II; erythropoietin (EPO); osteoinductive factors; interferons such as interferon-alpha, -beta and -gamma colony stimulating factors (CSFs) such as macrophage-CSF (M-CSF);  
5 granulocyte-macrophage-CSF (GM-CSF); and granulocyte-CSF (G-CSF); interleukins (ILs) such as IL-1, IL-1alpha, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-11, IL-12; a tumor necrosis factor such as TNF-alpha or TNF-beta; and other polypeptide factors including LIF and  
10 kit ligand (KL). As used herein, the term cytokine includes proteins from natural sources or from recombinant cell culture and biologically active equivalents of the native sequence cytokines.

The terms "pre-treatment" or "pre-treat" as used herein refers to exposure of the mammalian cell(s) to CPT-11 (or other chemotherapeutic agent) prior to exposure to Apo-2L receptor agonist(s). The pre-treatment of mammalian cells, particularly, cancer cells, with CPT-11 is believed to sensitize the cancer cells to Apo-2 ligand receptor agonist by enhancing or up-regulating expression of DR4 or DR5 receptor(s) in or on said cancer cells. Preferably, the amount of CPT-11 employed to pre-treat the cells will be an amount sufficient to enhance or up-regulate  
15 expression of DR4 or DR5 receptor(s) in or on said mammalian cells by about 0.5 to about 5-fold, more preferably, by about 1 to about 4-fold, and more preferably, by about 2 to about 4-fold, as compared to the same mammalian cells which are not exposed to CPT-11 under the same  
20 conditions.

"Treatment" or "therapy" refer to both therapeutic treatment and prophylactic or preventative measures.

The term "effective amount" refers to an amount of a drug effective to treat a disease or disorder in a mammal. In the case of cancer, the  
30 therapeutically effective amount of the drug may reduce the number of cancer cells; reduce the tumor size; inhibit (i.e., slow to some extent and preferably stop) cancer cell infiltration into peripheral organs; inhibit (i.e., slow to some extent and preferably stop) tumor metastasis; inhibit, to some extent, tumor growth; and/or relieve to some extent one  
35 or more of the symptoms associated with the disorder. To the extent the drug may prevent growth and/or kill existing cancer cells, it may be cytostatic and/or cytotoxic. For cancer therapy, efficacy in vivo can, for example, be measured by assessing tumor burden or volume, the time to disease progression (TTP) and/or determining the response rates (RR).

40 "Mammal" for purposes of treatment or therapy refers to any animal classified as a mammal, including humans, domestic and farm animals, and zoo, sports, or pet animals, such as dogs, horses, cats, cows, etc. Preferably, the mammal is human.

The terms "cancer", "cancerous", or "malignant" refer to or describe  
45 the physiological condition in mammals that is typically characterized by

unregulated cell growth. Examples of cancer include but are not limited to, carcinoma, lymphoma, blastoma, sarcoma, and leukemia. More particular examples of such cancers include colon cancer, colorectal cancer, rectal cancer, squamous cell cancer, small-cell lung cancer, non-small cell lung cancer, Hodgkin's and non-Hodgkin's lymphoma, testicular cancer, myeloma, esophageal cancer, gastrointestinal cancer, renal cancer, pancreatic cancer, glioblastoma, cervical cancer, ovarian cancer, glioma, liver cancer, bladder cancer, hepatoma, breast cancer, endometrial carcinoma, salivary gland carcinoma, kidney cancer, liver cancer, prostate cancer, vulval cancer, thyroid cancer, hepatic carcinoma and various types of head and neck cancer.

## II. Methods and Materials

### A. METHODS

Generally, the methods of the invention for inducing apoptosis in mammalian cells comprise exposing the cells to an effective amount of Apo-2 ligand and CPT-11 or an effective amount of Apo-2L receptor agonist antibody and CPT-11, wherein said cells are exposed to the CPT-11 prior to being exposed to said Apo-2L or Apo-2L receptor agonist antibody. Optionally, the amount of Apo-2L (or agonist antibody) employed will be an amount effective to induce apoptosis. Optionally, the amount of CPT-11 employed will be an amount effective to enhance expression of DR4 or DR5 receptor(s) in or on said cells. This can be accomplished in vivo or ex vivo in accordance, for instance, with the methods described below and in the Examples. Exemplary conditions or disorders to be treated with the Apo-2 ligand or agonist antibody and CPT-11 include benign or malignant cancer.

#### 1. Elements of Apoptotic Machinery

A further understanding of certain elements of the apoptotic machinery that correlate with an increase in killing activity can facilitate the practice of methods for inducing apoptosis in mammalian cells. In this context, the data provided in Example 3 identifies elements in the apoptotic machinery that correlate with increased killing activity of Apo2L/TRAIL plus CPT-11 treatment.

As discussed in detail below, Apo2L/TRAIL treatment of responsive tumor cells, but not normal cells, can induce a transient upregulation of DR5 receptor(s). Likewise, CPT-11 exposure can result in upregulation of DR5 receptor(s) and/or DR4 receptor(s). The combined Apo2L/TRAIL plus CPT-11 treatment for about 24 hours also resulted in augmented expression of DR4 and DR5 in comparison to controls. Moreover, the pre-treatment of various cells with CPT-11 for 20-22 hours followed by two hours with Apo2L/TRAIL produced the highest induction of DR5 and DR4 mRNA, as well as caspase -3- like cleavage/activation and apoptosis. The addition of the caspase inhibitor Z-VAD was found to further intensify DR5 and DR4 mRNA expression levels.

The data provided in Example 3 provides evidence that CPT-11 and Apo2L/TRAIL induce apoptosis by distinct, p53-dependent and p53 independent pathways, respectively. Specifically, Apo2L/TRAIL treatment of HCT-116 cells alone did not induce p53 expression, while CPT-11 and CPT-11 in combination with Apo2L/TRAIL resulted in strong induction of p53 protein. In addition, Apo2L/TRAIL mediated a transient upregulation of DR5 mRNA expression, while CPT-11 increased both DR5 and DR4 mRNA expression. CPT-11 alone induced a substantial upregulation of p21 protein, a p53 inducible, cyclin-dependent kinase inhibitor, that has been implicated in cell cycle arrest. CPT-11-induced accumulation of p21 was prevented by co-treatment with Apo2L/TRAIL in a caspase dependent fashion. Furthermore rather than accumulation at G2-M phase, cells co-treated with Apo2L/TRAIL underwent apoptosis. Thus, combined Apo2L/TRAIL and CPT-11 treatment led to degradation of p21 and to upregulation of DR4 and DR5, directing cancer cells towards an apoptotic pathway rather than cell cycle arrest and possible DNA repair. This is in clear agreement with the enhanced anti-tumor activity shown *in vivo* with the combination treatment (Ashkenazi, et. al. J. Clinical Investigation, 104: 155-162 (1999); Gliniak et al., Cancer Research, 59:6153-6158 (1999)) and these data provide a potential mechanism by which Apo2L/TRAIL and CPT-11 treatment mediates enhanced anti-tumor activity.

The data presented in Example 3 show no changes in FLICE inhibitory protein (FLIP) protein expression in HCT116 cells undergoing apoptosis, ruling out a significant anti-apoptotic involvement for FLIP in this experimental system, data which is in agreement with studies in melanoma tumors (Griffith et al., Current Opinion in Immunology, 10: 559-563 (1998); Leverkus et al., Cancer Research, 60:553-559 (2000); Zhang et al., Cancer Research, 59:2747-2753 (1999)). In addition, the presence of the general caspase inhibitor, ZVAD, effectively blocked Apo2L/TRAIL-mediated apoptosis, degradation of p21, and disruption of the G2-M phase cell arrest mediated by CPT-11 which provides evidence that p21 plays a regulatory role in Apo2L/TRAIL-mediated apoptosis (see also Xu et al., Biochem. Biophys. Res. Comm., 269: 179-190 (2000)). However, induction of p21 overexpression in these conditions prevented apoptosis by inhibition of proximal caspase activation.

The data presented herein describes a novel mechanism by which CPT-11 and Apo2L/TRAIL, two agents that mediate apoptosis through distinct pathways, DNA damage and death signaling receptors, respectively, can act in concert. Namely, Apo2L/TRAIL inhibition of p21-induction by CPT-11 can preclude accumulation of cells in G2/M cell cycle arrest, and therefore promotes increased apoptosis. In addition, the upregulation of death receptors by the combination of these agents may also contribute to the observed enhanced apoptotic activity.

## 2. Modulating Apo-2L receptor agonist induced Apoptosis

As disclosed herein, it is possible to modulate and augment the apoptosis in mammalian cancer cells which occurs when cells are exposed to an effective amount of CPT-11 and an Apo-2L receptor agonist by administering the CPT-11 prior to the administration of the Apo-2 ligand receptor agonist. Specifically, as shown in Example 3 and Figure 5, the pre-treatment of cells with CPT-11 for 20-22 hours followed by two hours with Apo2L/TRAIL produced the highest induction of DR5 and DR4 mRNA, as well as caspase -3- like cleavage/activation and apoptosis. Therefore, an important aspect of the invention are improved methods of using Apo-2L receptor agonists and a chemotherapeutic agent such as CPT-11 to induce apoptosis in mammalian cells, wherein the methods comprise pre-treating the cells with the chemotherapeutic agent prior to their treatment with the Apo-2L receptor agonist.

Methods of pre-treating mammalian cells with a chemotherapeutic agent such as CPT-11 prior to their treatment with the Apo-2L receptor agonist(s) can have a number of advantages over the simultaneous administration of these agents. In particular, as noted above, these methods can facilitate treatment modalities by identifying the optimal conditions for the combined administration of these agents. Consequently, by identifying methods to optimize an apoptotic response, medical practitioners may be able to dispense these agents in a more convenient and patient friendly format. Specifically, employing methods which optimize an apoptotic response, medical practitioners may administer these agents in a single bolus rather than in multiple injections, administer lower concentrations of these agents or administer these agents for shorter periods of time.

Additional chemotherapeutic agents having physiological effects that are similar to those of CPT-11 can also be used in the methods disclosed herein. Specifically, exposure to different anti-cancer genotoxic-stress chemicals such as doxorubicin, etoposide, CDDP and gamma irradiation treatments can also result in selective p53-dependent upregulation of the Apo2/TRAIL death-receptor DR5 in a number of tumor cell lines (see e.g. Kim et al., Clin. Cancer Res. 6(2): 335-346 (2000); Gibson et al., Mol. Cell Biol. 20(1): 205-212 (2000); Keane et al., Cancer Res. 59(3): 734-741 (1999); Nagane et al., Cancer Res. 60(4): 847-853 (2000); Wu et al., Nature Genetics. 17:141-3 (1997) and Wu et al., Oncogene. 18: 6411-6418 (1999)). Upregulation of DR5 in a p53-independent fashion has also been demonstrated by treatment of tumor cells with TNF- $\alpha$  (Sheikh et al., Cancer Research 58: 1593-1598 (1998)) or by several chemotherapeutic agents in different human glioma cell lines (Nagane et al., Cancer Research 60:847-853 (2000)). Furthermore, upregulation of DR5 correlated in most cases with increased responsiveness to caspase-dependent Apo2L/TRAIL-mediated apoptosis (Chinnaiyan et al., Proc. Nat. Acad. Sci., 97:1754-1759 (2000)).

As disclosed herein one can enhance Apo-2L receptor agonist mediated apoptosis in mammalian cancer cells by pre-treating the cells with an agent that modulates the cellular apoptotic machinery associated with increased killing activity. Typical embodiments of the invention disclosed herein include a method for sensitizing cells to Apo-2L receptor agonist mediated apoptosis by pre-treating the cells with an agent that effects one or more physiological events including the upregulation of DR4, the upregulation of DR5 and/or the induction of p53 protein. Preferably the agent is selected from the group consisting of CPT-11, doxorubicin, 5-fluorouracil, interferon (e.g., interferon alpha or interferon gamma), etoposide, cis-diamminedichloroplatinum(II) (CDDP), TNF- $\alpha$  and gamma irradiation. In highly preferred embodiments, the agent is CPT-11.

In accordance with one embodiment of the invention, there is provided a method of inducing apoptosis in mammalian cancer cells comprising exposing the cells to an effective amount of CPT-11 and an Apo-2 ligand receptor agonist, wherein the cells are exposed to CPT-11 prior to the Apo-2 ligand receptor agonist. Preferably, in these methods, the amount of administered CPT-11 results in an upregulation of DR4 and/or DR5 in or on said cells. The upregulation or enhanced expression of DR4 and/or DR5 may be assayed and measured, as compared to control cells not exposed to CPT-11, using known techniques such as by measuring expression of DR4 or DR5 mRNA, and including those techniques described in the Examples. Such assays may be conducted at selected time points following exposure of the cells to CPT-11 to determine the optimum desired time period for pre-treatment that may induce the desired or optimum upregulation of DR4 or DR5. Using *in vitro* assay methods, Applicants have found that induction of DR5 expression by CPT-11 can be observed after two hours exposure or incubation, and particularly, that DR5 expression can be induced *in vitro* following exposure of cells to 50 microgram/ml CPT-11 for 6 hours. Optionally, the cells may be exposed to the CPT-11 from about 1 hour to about 5 days, preferably about 2 hours to about 24, 48, or 72 hours, and more preferably about 6 hours to about 24 or 48 hours prior to exposure to Apo-2L receptor agonist(s). In the methods, the Apo-2 ligand receptor agonist typically comprises Apo2L/TRAIL or anti-DR4 receptor antibody. Additional embodiments of the invention include variations on these methods such as those that employ additional therapeutic modalities, such as exposing the cancer cells to one or more growth inhibitory agents or radiation. In preferred embodiments of the methods, the cancer cells comprise colorectal cancer cells.

Typically, an effective dose of CPT-11 is an amount sufficient to upregulate DR4 or DR5, or both DR4 and DR5, in or on the mammalian cells exposed to the CPT-11. In addition, an effective dose of CPT-11 typically induces p53 protein. Typical doses of CPT-11 employed include



standard clinical doses according to the Physician's Desk Reference (PDR) and may include a range from about 1 microgram/ml to about 100 microgram/ml, and optionally from about 2 microgram/ml to about 50 microgram/ml, and for clinical use, may preferably include a range from  
5 about 0.05 mg/kg to about 2.5 mg/kg, while typical doses of Apo-2 ligand may include a range from 0.1 mg/kg to about 12.0 mg/kg.

B. MATERIALS

The Apo-2L which can be employed in the methods includes the Apo-2L polypeptides described in Pitti et al., supra, WO 97/25428, supra, and  
10 WO97/01633, supra (the polypeptides referred to as TRAIL). It is contemplated that various forms of Apo-2L may be used, such as the full length polypeptide as well as soluble forms of Apo-2L which comprise an extracellular domain (ECD) sequence. Examples of such soluble ECD sequences include polypeptides comprising amino acids 114-281, 95-281,  
15 91-281 or 92-281 of the Apo-2L sequence shown in Figure 1A of Pitti et al., J. Biol. Chem., 271:12687-12690 (1996) and SEQ ID NO:1 herein. It is presently believed that the polypeptide comprising amino acids 92-281 is a naturally cleaved form of Apo-2L. Applicants have expressed human Apo-2L in CHO cells and found that the 92-281 polypeptide is the  
20 expressed form of Apo-2L. Modified forms of Apo-2L, such as the covalently modified forms described in WO 97/25428 are included. In particular, Apo-2L linked to a non-proteinaceous polymer such as polyethylene glycol is included for use in the present methods. The Apo-2L polypeptide can be made according to any of the methods described in  
25 WO 97/25428.

Variants of Apo-2 ligand having apoptotic activity which can be used in the methods include, for example, those identified by alanine scanning techniques. Particular substitutional variants comprise amino acids 91-281 of Figure 1A of Pitti et al., J. Biol. Chem., 271:12687-  
30 12690 (1996) in which at least one of the amino acids at positions 203, 218 or 269 are substituted by an alanine residue. Optionally, the Apo-2 ligand variants may include one or more of these three different site substitutions.

It is contemplated that a molecule which mimics the apoptotic  
35 activity of Apo-2L may alternatively be employed in the presently disclosed methods. Examples of such molecules include agonistic antibodies which can induce apoptosis in at least a comparable or like manner to Apo-2L. In particular, these agonist antibodies would comprise antibodies which bind one or more of the receptors for Apo-2L.  
40 Preferably, the agonist antibody is directed to an Apo-2L receptor which includes a cytoplasmic death domain, such as DR4 or DR5. Even more preferably, the agonist antibody binds to such a receptor and binding can be determined, e.g., using FACS analysis or ELISA, such as described in Example 2. Agonist antibodies directed to the receptor called DR5 (or  
45 Apo-2) have been prepared using fusion techniques such as described

below. One of the DR5 or Apo-2 receptor agonist antibodies is referred to as 3F11.39.7 and has been deposited with ATCC as deposit no. HB-12456 on January 13, 1998. Other DR5 receptor antibodies include 3H3.14.5, deposited with ATCC as shown herein. Agonist activity of the Apo-2L receptor antibodies can be determined using various methods for assaying for apoptotic activity, and optionally, apoptotic activity of such antibody can be determined by assaying the antibody, alone or in a cross-linked form using Fc immunoglobulin or complement (described below), in the assay described in Example 2 to examine apoptosis of 9D cells or other cells expressing an Apo-2L receptor such as DR4 or DR5.

Additionally, agonist antibodies directed to another Apo-2L receptor, called DR4, have also been prepared. One of the DR4 agonist antibodies is referred to as 4H6.17.8 and has been deposited with ATCC as deposit no. HB-12455 on January 13, 1998. Still further agonist DR4 antibodies include the antibodies 4E7.24.3, 1H5.25.9, 4G7.18.8, and 5G11.17.1 which have been deposited with ATCC, as shown below. Agonist activity of the Apo-2L receptor antibodies can be determined using various methods for assaying for apoptotic activity, and optionally, apoptotic activity of such antibody can be determined by assaying the antibody, alone or in a cross-linked form using Fc immunoglobulin or complement (described below), in the assay described in Example 2 to examine apoptosis of 9D cells or other cells expressing an Apo-2L receptor such as DR4 or DR5.

Agonist antibodies contemplated by the invention include antibodies which bind a single Apo-2L receptor or more than one Apo-2L receptor. An antibody which binds more than one Apo-2L receptor can be characterized as an antibody that "cross-reacts" with two or more different antigens and capable of binding to each of the different antigens, e.g. as determined by ELISA or FACS as in the examples below. Optionally, an antibody which "specifically cross-reacts" with two or more different antigens is one which binds to a first antigen and further binds to a second different antigen, wherein the binding ability of the antibody for the second antigen at an antibody concentration of about 10 $\mu$ g/mL is from about 50% to about 100% (preferably from about 75% to about 100%) of the binding ability of the first antigen as determined in a capture ELISA (such as in the examples below). For example, the antibody may bind specifically to DR5 (the "first antigen") and specifically cross-react with another Apo-2L receptor such as DR4 (the "second antigen"), wherein the extent of binding of about 10 $\mu$ g/mL of the antibody to DR4 is about 50% to about 100% of the binding ability of the antibody for DR5 in the capture ELISA herein. Various cross-reactive antibodies to Apo-2L receptors are described in further detail in International Patent application number PCT/US99/13197.

As described below, exemplary forms of such antibodies include polyclonal, monoclonal, humanized, bispecific, and heteroconjugate antibodies.

### 1. Polyclonal Antibodies

5       The antibodies of the invention may comprise polyclonal antibodies. Methods of preparing polyclonal antibodies are known to the skilled artisan. Polyclonal antibodies can be raised in a mammal, for example, by one or more injections of an immunizing agent and, if desired, an adjuvant. Typically, the immunizing agent and/or adjuvant will be injected  
10       in the mammal by multiple subcutaneous or intraperitoneal injections. The immunizing agent may include a DR4 or DR5 polypeptide (or a DR4 or DR5 ECD) or a fusion protein thereof. It may be useful to conjugate the immunizing agent to a protein known to be immunogenic in the mammal being immunized. Examples of such immunogenic proteins include but are not  
15       limited to keyhole limpet hemocyanin, serum albumin, bovine thyroglobulin, and soybean trypsin inhibitor. Examples of adjuvants which may be employed include Freund's complete adjuvant and MPL-TDM adjuvant (monophosphoryl Lipid A, synthetic trehalose dicorynomycolate). The immunization protocol may be selected by one skilled in the art without  
20       undue experimentation. The mammal can then be bled, and the serum assayed for antibody titer. If desired, the mammal can be boosted until the antibody titer increases or plateaus.

### 2. Monoclonal Antibodies

      The antibodies of the invention may, alternatively, be monoclonal  
25       antibodies. Monoclonal antibodies may be prepared using hybridoma methods, such as those described by Kohler and Milstein, Nature, 256:495 (1975). In a hybridoma method, a mouse, hamster, or other appropriate host animal, is typically immunized with an immunizing agent to elicit lymphocytes that produce or are capable of producing antibodies that will  
30       specifically bind to the immunizing agent. Alternatively, the lymphocytes may be immunized *in vitro*.

      The immunizing agent will typically include a DR4 or DR5 polypeptide or a fusion protein thereof, such as a DR4 or DR5 ECD-IgG fusion protein.

      Generally, either peripheral blood lymphocytes ("PBLs") are used if  
35       cells of human origin are desired, or spleen cells or lymph node cells are used if non-human mammalian sources are desired. The lymphocytes are then fused with an immortalized cell line using a suitable fusing agent, such as polyethylene glycol, to form a hybridoma cell [Goding, Monoclonal Antibodies: Principles and Practice, Academic Press, (1986) pp. 59-103].

40       Immortalized cell lines are usually transformed mammalian cells, particularly myeloma cells of rodent, bovine and human origin. Usually, rat or mouse myeloma cell lines are employed. The hybridoma cells may be cultured in a suitable culture medium that preferably contains one or more substances that inhibit the growth or survival of the unfused,  
45       immortalized cells. For example, if the parental cells lack the enzyme

hypoxanthine guanine phosphoribosyl transferase (HGPRT or HPRT), the culture medium for the hybridomas typically will include hypoxanthine, aminopterin, and thymidine ("HAT medium"), which substances prevent the growth of HGPRT-deficient cells.

5 Preferred immortalized cell lines are those that fuse efficiently, support stable high level expression of antibody by the selected antibody-producing cells, and are sensitive to a medium such as HAT medium. More preferred immortalized cell lines are murine myeloma lines, which can be obtained, for instance, from the Salk Institute Cell Distribution Center,  
10 San Diego, California and the American Type Culture Collection, Manassas, Virginia. An example of such a murine myeloma cell line is P3X63AgU.1 described in Example 2 below. Human myeloma and mouse-human heteromyeloma cell lines also have been described for the production of human monoclonal antibodies [Kozbor, J. Immunol., 133:3001 (1984); Brodeur et al.,  
15 Monoclonal Antibody Production Techniques and Applications, Marcel Dekker, Inc., New York, (1987) pp. 51-63].

The culture medium in which the hybridoma cells are cultured can then be assayed for the presence of monoclonal antibodies directed against the Apo-2L receptor. Preferably, the binding specificity of monoclonal  
20 antibodies produced by the hybridoma cells is determined by immunoprecipitation or by an in vitro binding assay, such as radioimmunoassay (RIA) or enzyme-linked immunoabsorbent assay (ELISA). Such techniques and assays are known in the art. The binding affinity of the monoclonal antibody can, for example, be determined by the Scatchard analysis of Munson and Pollard, Anal. Biochem., 107:220 (1980).  
25

After the desired hybridoma cells are identified, the clones may be subcloned by limiting dilution procedures and grown by standard methods [Goding, supra]. Suitable culture media for this purpose include, for example, Dulbecco's Modified Eagle's Medium or RPMI-1640 medium.  
30 Alternatively, the hybridoma cells may be grown in vivo as ascites in a mammal.

The monoclonal antibodies secreted by the subclones may be isolated or purified from the culture medium or ascites fluid by conventional immunoglobulin purification procedures such as, for example, protein A-  
35 Sepharose, hydroxylapatite chromatography, gel electrophoresis, dialysis, or affinity chromatography.

The monoclonal antibodies may also be made by recombinant DNA methods, such as those described in U.S. Patent No. 4,816,567. DNA encoding the monoclonal antibodies of the invention can be readily  
40 isolated and sequenced using conventional procedures (e.g., by using oligonucleotide probes that are capable of binding specifically to genes encoding the heavy and light chains of murine antibodies). The hybridoma cells of the invention serve as a preferred source of such DNA. Once isolated, the DNA may be placed into expression vectors, which are then  
45 transfected into host cells such as simian COS cells, Chinese hamster

ovary (CHO) cells, or myeloma cells that do not otherwise produce immunoglobulin protein, to obtain the synthesis of monoclonal antibodies in the recombinant host cells. The DNA also may be modified, for example, by substituting the coding sequence for human heavy and light chain constant domains in place of the homologous murine sequences [U.S. Patent No. 4,816,567; Morrison et al., supra] or by covalently joining to the immunoglobulin coding sequence all or part of the coding sequence for a non-immunoglobulin polypeptide. Such a non-immunoglobulin polypeptide can be substituted for the constant domains of an antibody of the invention, or can be substituted for the variable domains of one antigen-combining site of an antibody of the invention to create a chimeric bivalent antibody. Optionally, chimeric antibodies can be constructed which include at least one variable or hypervariable domain of an anti-Apo-2L receptor antibody selected from the 4H6.17.8, 3F11.39.7, 4E7.24.3, 1H5.25.9, 4G7.18.8, 5G11.17.1, and 3H3.14.5 antibodies disclosed herein.

Optionally, the agonist antibodies of the present invention will bind to the same epitope(s) as any of the 4H6.17.8, 3F11.39.7, 4E7.24.3, 1H5.25.9, 4G7.18.8, 5G11.17.1, and 3H3.14.5 antibodies disclosed herein. This can be determined by conducting various assays, such as described herein. For instance, to determine whether a monoclonal antibody has the same specificity as the DR4 or DR5 antibodies specifically referred to herein, one can compare its activity in blocking assays or apoptosis induction assays.

The antibodies of the invention include "cross-linked" antibodies. The term "cross-linked" as used herein refers to binding of at least two IgG molecules together to form one (or single) molecule. The Apo-2L receptor antibodies may be cross-linked using various linker molecules, optionally the DR4 antibodies are cross-linked using an anti-IgG molecule, complement, chemical modification or molecular engineering. It is appreciated by those skilled in the art that complement has a relatively high affinity to antibody molecules once the antibodies bind to cell surface membrane. Accordingly, it is believed that complement may be used as a cross-linking molecule to link two or more antibodies bound to cell surface membrane. Among the various murine Ig isotypes, IgM, IgG2a and IgG2b are known to fix complement.

The antibodies of the invention may optionally comprise dimeric antibodies, as well as multivalent forms of antibodies. Those skilled in the art may construct such dimers or multivalent forms by techniques known in the art and using the anti-Apo-2L receptor antibodies herein.

The antibodies of the invention may also comprise monovalent antibodies. Methods for preparing monovalent antibodies are well known in the art. For example, one method involves recombinant expression of immunoglobulin light chain and modified heavy chain. The heavy chain is truncated generally at any point in the Fc region so as to prevent heavy chain crosslinking. Alternatively, the relevant cysteine residues are

substituted with another amino acid residue or are deleted so as to prevent crosslinking.

In vitro methods are also suitable for preparing monovalent antibodies. Digestion of antibodies to produce fragments thereof, particularly, Fab fragments, can be accomplished using routine techniques known in the art. For instance, digestion can be performed using papain. Examples of papain digestion are described in WO 94/29348 published 12/22/94 and U.S. Patent No. 4,342,566. Papain digestion of antibodies typically produces two identical antigen binding fragments, called Fab fragments, each with a single antigen binding site, and a residual Fc fragment. Pepsin treatment yields an  $F(ab')_2$  fragment that has two antigen combining sites and is still capable of cross-linking antigen.

The Fab fragments produced in the antibody digestion also contain the constant domains of the light chain and the first constant domain ( $CH_1$ ) of the heavy chain. Fab' fragments differ from Fab fragments by the addition of a few residues at the carboxy terminus of the heavy chain  $CH_1$  domain including one or more cysteines from the antibody hinge region. Fab'-SH is the designation herein for Fab' in which the cysteine residue(s) of the constant domains bear a free thiol group.  $F(ab')_2$  antibody fragments originally were produced as pairs of Fab' fragments which have hinge cysteines between them. Other chemical couplings of antibody fragments are also known.

Single chain Fv fragments may also be produced, such as described in Iliades et al., FEBS Letters, 409:437-441 (1997). Coupling of such single chain fragments using various linkers is described in Kortt et al., Protein Engineering, 10:423-433 (1997).

In addition to the antibodies described above, it is contemplated that chimeric or hybrid antibodies may be prepared in vitro using known methods in synthetic protein chemistry, including those involving crosslinking agents. For example, immunotoxins may be constructed using a disulfide exchange reaction or by forming a thioether bond. Examples of suitable reagents for this purpose include iminothiolate and methyl-4-mercaptopbutyrimidate.

The Apo-2L receptor antibodies of the invention may further comprise humanized antibodies or human antibodies. Humanized forms of non-human (e.g., murine) antibodies are chimeric immunoglobulins, immunoglobulin chains or fragments thereof (such as Fv, Fab, Fab',  $F(ab')_2$  or other antigen-binding subsequences of antibodies) which contain minimal sequence derived from non-human immunoglobulin. Humanized antibodies include human immunoglobulins (recipient antibody) in which residues from a complementary determining region (CDR) of the recipient are replaced by residues from a CDR of a non-human species (donor antibody) such as mouse, rat or rabbit having the desired specificity, affinity and capacity. In some instances, Fv framework residues of the human immunoglobulin are

replaced by corresponding non-human residues. Humanized antibodies may also comprise residues which are found neither in the recipient antibody nor in the imported CDR or framework sequences. In general, the humanized antibody will comprise substantially all of at least one, and typically two, variable domains, in which all or substantially all of the CDR regions correspond to those of a non-human immunoglobulin and all or substantially all of the FR regions are those of a human immunoglobulin consensus sequence. The humanized antibody optimally also will comprise at least a portion of an immunoglobulin constant region (Fc), typically that of a human immunoglobulin [Jones et al., Nature, 321:522-525 (1986); Riechmann et al., Nature, 332:323-329 (1988); and Presta, Curr. Op. Struct. Biol., 2:593-596 (1992)].

Methods for humanizing non-human antibodies are well known in the art. Generally, a humanized antibody has one or more amino acid residues introduced into it from a source which is non-human. These non-human amino acid residues are often referred to as "import" residues, which are typically taken from an "import" variable domain. Humanization can be essentially performed following the method of Winter and co-workers [Jones et al., Nature, 321:522-525 (1986); Riechmann et al., Nature, 332:323-327 (1988); Verhoeven et al., Science, 239:1534-1536 (1988)], by substituting rodent CDRs or CDR sequences for the corresponding sequences of a human antibody. Accordingly, such "humanized" antibodies are chimeric antibodies (U.S. Patent No. 4,816,567), wherein substantially less than an intact human variable domain has been substituted by the corresponding sequence from a non-human species. In practice, humanized antibodies are typically human antibodies in which some CDR residues and possibly some FR residues are substituted by residues from analogous sites in rodent antibodies. Sources of such import residues or import variable domains (or CDRs) include the deposited anti-Apo-2L receptor antibodies 4H6.17.8, 3F11.39.7, 4E7.24.3, 1H5.25.9, 4G7.18.8, 5G11.17.1, and 3H3.14.5 disclosed herein.

The choice of human variable domains, both light and heavy, to be used in making the humanized antibodies is very important in order to reduce antigenicity. According to the "best-fit" method, the sequence of the variable domain of a rodent antibody is screened against the entire library of known human variable domain sequences. The human sequence which is closest to that of the rodent is then accepted as the human framework (FR) for the humanized antibody [Sims et al., J. Immunol., 151:2296-2308 (1993); Chothia and Lesk, J. Mol. Biol., 196:901-917 (1987)]. Another method uses a particular framework derived from the consensus sequence of all human antibodies of a particular subgroup of light or heavy chains. The same framework may be used for several different humanized antibodies [Carter et al., Proc. Natl. Acad. Sci. USA, 89:4285-4289 (1992); Presta et al., J. Immunol., 151:2623-2632 (1993)].

It is further important that antibodies be humanized with retention of high affinity for the antigen and other favorable biological properties. To achieve this goal, according to a preferred method, humanized antibodies are prepared by a process of analysis of the parental sequences and various conceptual humanized products using three dimensional models of the parental and humanized sequences. Three dimensional immunoglobulin models are commonly available and are familiar to those skilled in the art. Computer programs are available which illustrate and display probable three-dimensional conformational structures of selected candidate immunoglobulin sequences. Inspection of these displays permits analysis of the likely role of the residues in the functioning of the candidate immunoglobulin sequence, i.e., the analysis of residues that influence the ability of the candidate immunoglobulin to bind its antigen. In this way, FR residues can be selected and combined from the consensus and import sequence so that the desired antibody characteristic, such as increased affinity for the target antigen(s), is achieved. In general, the CDR residues are directly and most substantially involved in influencing antigen binding [see, WO 94/04679 published 3 March 1994].

Human monoclonal antibodies may be made via an adaptation of the hybridoma method first described by Kohler and Milstein by using human B lymphocytes as the fusion partner. Human B lymphocytes producing an antibody of interest may, for example, be isolated from a human individual, after obtaining informed consent. For instance, the individual may be producing antibodies against an autoantigen as occurs with certain disorders such as systemic lupus erythematosus (Shoenfeld et al. J. Clin. Invest., 70:205 (1982)), immune-mediated thrombocytopenic purpura (ITP) (Nugent et al. Blood, 70(1): 16-22 (1987)), or cancer. Alternatively, or additionally, lymphocytes may be immunized *in vitro*. For instance, one may expose isolated human peripheral blood lymphocytes *in vitro* to a lysomotropic agent (e.g. L-leucine-O-methyl ester, L-glutamic acid dimethyl ester or L-leucyl-L-leucine-O-methyl ester) (US Patent No. 5,567,610, Borrebaeck et al.); and/or T-cell depleted human peripheral blood lymphocytes may be treated *in vitro* with adjuvants such as 8-mercaptoguanosine and cytokines (US Patent No. 5,229,275, Goroff et al.).

The B lymphocytes recovered from the subject or immunized *in vitro*, are then generally immortalized in order to generate a human monoclonal antibody. Techniques for immortalizing the B lymphocyte include, but are not limited to: (a) fusion of the human B lymphocyte with human, murine myelomas or mouse-human heteromyeloma cells; (b) viral transformation (e.g. with an Epstein-Barr virus; see Nugent et al., *supra*, for example); (c) fusion with a lymphoblastoid cell line; or (d) fusion with lymphoma cells.



Lymphocytes may be fused with myeloma cells using a suitable fusing agent, such as polyethylene glycol, to form a hybridoma cell (Goding, Monoclonal Antibodies: Principles and Practice, pp.59-103 (Academic Press, 1986)). The hybridoma cells thus prepared are seeded and grown in  
5 a suitable culture medium that preferably contains one or more substances that inhibit the growth or survival of the unfused, parental myeloma cells. For example, if the parental myeloma cells lack the enzyme hypoxanthine guanine phosphoribosyl transferase (HGPRT or HPRT), the culture medium for the hybridomas typically will include hypoxanthine,  
10 aminopterin, and thymidine (HAT medium), which substances prevent the growth of HGPRT-deficient cells. Suitable human myeloma and mouse-human heteromyeloma cell lines have been described (Kozbor, *J. Immunol.*, 133:3001 (1984); Brodeur et al., Monoclonal Antibody Production Techniques and Applications, pp. 51-63 (Marcel Dekker, Inc., New York,  
15 1987)). Culture medium in which hybridoma cells are growing is assayed for production of monoclonal antibodies directed against the antigen. Preferably, the binding specificity of monoclonal antibodies produced by hybridoma cells is determined by immunoprecipitation or by an *in vitro* binding assay, such as radioimmunoassay (RIA) or enzyme-linked  
20 immunoabsorbent assay (ELISA).

After hybridoma cells are identified that produce antibodies of the desired specificity, affinity, and/or activity, the clones may be subcloned by limiting dilution procedures and grown by standard methods (Goding, Monoclonal Antibodies: Principles and Practice, pp.59-103  
25 (Academic Press, 1986)). Suitable culture media for this purpose include, for example, D-MEM or RPMI-1640 medium. The monoclonal antibodies secreted by the subclones are suitably separated from the culture medium, ascites fluid, or serum by conventional immunoglobulin purification procedures such as, for example, protein A chromatography,  
30 gel electrophoresis, dialysis, or affinity chromatography.

Human antibodies may also be generated using a non-human host, such as a mouse, which is capable of producing human antibodies. As noted above, transgenic mice are now available that are capable, upon immunization, of producing a full repertoire of human antibodies in the  
35 absence of endogenous immunoglobulin production. For example, it has been described that the homozygous deletion of the antibody heavy-chain joining region (J<sub>H</sub>) gene in chimeric and germ-line mutant mice results in complete inhibition of endogenous antibody production. Transfer of the human germ-line immunoglobulin gene array in such germ-line mutant mice  
40 will result in the production of human antibodies upon antigen challenge. See, e.g., Jakobovits et al., *Proc. Natl. Acad. Sci. USA*, 90:2551 (1993); Jakobovits et al., *Nature*, 362:255-258 (1993); Bruggermann et al., *Year in Immuno.*, 7:33 (1993); US Patent No. 5,591,669; US Patent No. 5,589,369; and US Patent No. 5,545,807. Human antibodies may also be  
45 prepared using SCID-hu mice (Duchosal et al. *Nature* 355:258-262 (1992)).

In another embodiment, the human antibody may be selected from a human antibody phage display library. The preparation of libraries of antibodies or fragments thereof is well known in the art and any of the known methods may be used to construct a family of transformation vectors which may be introduced into host cells. Libraries of antibody light and heavy chains in phage (Huse et al., *Science*, 246:1275 (1989)) or of fusion proteins in phage or phagemid can be prepared according to known procedures. See, for example, Vaughan et al., *Nature Biotechnology* 14:309-314 (1996); Barbas et al., *Proc. Natl. Acad. Sci., USA*, 88:7978-7982 (1991); Marks et al., *J. Mol. Biol.*, 222:581-597 (1991); Hoogenboom and Winter, *J. Mol. Biol.*, 227:381-388 (1992); Barbas et al., *Proc. Natl. Acad. Sci., USA*, 89:4457-4461 (1992); Griffiths et al., *EMBO Journal*, 13:3245-3260 (1994); de Kruif et al., *J. Mol. Biol.*, 248:97-105 (1995); WO 98/05344; WO 98/15833; WO 97/47314; WO 97/44491; WO 97/35196; WO 95/34648; US Patent No. 5,712,089; US Patent No. 5,702,892; US Patent No. 5,427,908; US Patent No. 5,403,484; US Patent No. 5,432,018; US Patent No. 5,270,170; WO 92/06176; WO 99/06587; US Patent No. 5,514,548; WO97/08320; and US Patent No. 5,702,892. The antigen of interest is panned against the phage library using procedures known in the field for selecting phage-antibodies which bind to the target antigen.

The Apo-2L receptor antibodies, as described herein, will optionally possess one or more desired biological activities or properties. Such antibodies may include but are not limited to chimeric, humanized, human, and affinity matured antibodies. As described above, the antibodies may be constructed or engineered using various techniques to achieve these desired activities or properties. In one embodiment, the Apo-2L receptor antibody will have a DR4 or DR5 receptor binding affinity of at least  $10^5 \text{ M}^{-1}$ , preferably at least in the range of  $10^6 \text{ M}^{-1}$  to  $10^7 \text{ M}^{-1}$ , more preferably, at least in the range of  $10^8 \text{ M}^{-1}$  to  $10^{12} \text{ M}^{-1}$  and even more preferably, at least in the range of  $10^9 \text{ M}^{-1}$  to  $10^{12} \text{ M}^{-1}$ . The binding affinity of the antibody can be determined without undue experimentation by testing the antibody in accordance with techniques known in the art, including Scatchard analysis (see Munson et al., *supra*). For example, a DR4 antibody can be assayed for binding affinity to the DR4-IgG receptor construct, as described in Example 2.

In another embodiment, the Apo-2L receptor antibody of the invention may bind the same epitope on DR4 or DR5 to which Apo-2L binds, or bind an epitope on DR4 or DR5 which coincides or overlaps with the epitope on DR4 or DR5, respectively, to which Apo-2L binds. The antibody may also interact in such a way to create a steric conformation which prevents Apo-2 ligand binding to DR4 or DR5. The epitope binding property of the antibody of the present invention may be determined using techniques known in the art. For instance, the antibody may be tested in an *in vitro* assay, such as a competitive inhibition assay, to determine

the ability of the antibody to block or inhibit binding of Apo-2L to DR4 or DR5. Optionally, the antibody may be tested in a competitive inhibition assay to determine the ability of, e.g., a DR4 antibody to inhibit binding of an Apo-2L polypeptide (such as described in Example 1) to a DR4-IgG construct (such as described in Example 2) or to a cell expressing DR4. Optionally, the antibody will be capable of blocking or inhibiting binding of Apo-2L to the receptor by at least 50%, preferably by at least 75% and even more preferably by at least 90%, which may be determined, by way of example, in an *in vitro* competitive inhibition assay using a soluble form of Apo-2 ligand (TRAIL) and a DR4 ECD-IgG (such as described in Example 2).

In a preferred embodiment, the antibody will comprise an agonist antibody having activity which mimics or is comparable to Apo-2 ligand (TRAIL). Preferably, such an agonistic DR4 or DR5 antibody will induce apoptosis in at least one type of cancer or tumor cell line or primary tumor. The apoptotic activity of an agonistic DR4 or DR5 antibody may be determined using known *in vitro* or *in vivo* assays. Examples of such *in vitro* and *in vivo* assays are described in detail in the Examples section below. *In vitro*, apoptotic activity can be determined using known techniques such as Annexin V binding. *In vivo*, apoptotic activity may be determined, e.g., by measuring reduction in tumor burden or volume.

### 3. Bispecific Antibodies

Bispecific antibodies are monoclonal, preferably human or humanized, antibodies that have binding specificities for at least two different antigens. In the present case, one of the binding specificities is for an Apo-2L receptor, the other one is for any other antigen, and preferably for a cell-surface protein or receptor or receptor subunit.

Methods for making bispecific antibodies are known in the art. Traditionally, the recombinant production of bispecific antibodies is based on the co-expression of two immunoglobulin heavy-chain/light-chain pairs, where the two heavy chains have different specificities [Milstein and Cuello, Nature, 305:537-539 (1983)]. Because of the random assortment of immunoglobulin heavy and light chains, these hybridomas (quadromas) produce a potential mixture of ten different antibody molecules, of which only one has the correct bispecific structure. The purification of the correct molecule is usually accomplished by affinity chromatography steps. Similar procedures are disclosed in WO 93/08829, published 13 May 1993, and in Traunecker et al., EMBO J., 10:3655-3659 (1991).

Antibody variable domains with the desired binding specificities (antibody-antigen combining sites) can be fused to immunoglobulin constant domain sequences. The fusion preferably is with an immunoglobulin heavy-chain constant domain, comprising at least part of the hinge, CH2, and CH3 regions. It is preferred to have the first heavy-chain constant region (CH1) containing the site necessary for light-chain binding present in at

least one of the fusions. DNAs encoding the immunoglobulin heavy-chain fusions and, if desired, the immunoglobulin light chain, are inserted into separate expression vectors, and are co-transfected into a suitable host organism. For further details of generating bispecific antibodies see, 5 for example, Suresh et al., Methods in Enzymology, 121:210 (1986).

#### 4. Heteroconjugate Antibodies

Heteroconjugate antibodies are also within the scope of the present invention. Heteroconjugate antibodies are composed of two covalently joined antibodies. Such antibodies have, for example, been proposed to 10 target immune system cells to unwanted cells [U.S. Patent No. 4,676,980], and for treatment of HIV infection [WO 91/00360; WO 92/200373; EP 03089]. It is contemplated that the antibodies may be prepared in vitro using known methods in synthetic protein chemistry, including those involving crosslinking agents. For example, immunotoxins may be constructed using a 15 disulfide exchange reaction or by forming a thioether bond. Examples of suitable reagents for this purpose include iminothiolate and methyl-4-mercaptobutyrimidate and those disclosed, for example, in U.S. Patent No. 4,676,980.

#### 5. Triabodies

20 Triabodies are also within the scope of the invention. Such antibodies are described for instance in Iliades et al., supra and Kortt et al., supra.

#### 6. Other Modifications

Other modifications of the Apo-2L receptor antibodies are 25 contemplated herein. The antibodies of the present invention may be modified by conjugating the antibody to a cytotoxic agent (like a toxin molecule) or a prodrug-activating enzyme which converts a prodrug (e.g. a peptidyl chemotherapeutic agent, see WO81/01145) to an active anti-cancer drug. See, for example, WO 88/07378 and U.S. Patent No. 4,975,278. This 30 technology is also referred to as "Antibody Dependent Enzyme Mediated Prodrug Therapy" (ADEPT).

The enzyme component of the immunoconjugate useful for ADEPT includes any enzyme capable of acting on a prodrug in such a way so as to covert it into its more active, cytotoxic form. Enzymes that are useful 35 in the method of this invention include, but are not limited to, alkaline phosphatase useful for converting phosphate-containing prodrugs into free drugs; arylsulfatase useful for converting sulfate-containing prodrugs into free drugs; cytosine deaminase useful for converting non-toxic 5-fluorocytosine into the anti-cancer drug, 5-fluorouracil; proteases, such 40 as serratia protease, thermolysin, subtilisin, carboxypeptidases and cathepsins (such as cathepsins B and L), that are useful for converting peptide-containing prodrugs into free drugs; caspases such as caspase-3; D-alanylcarboxypeptidases, useful for converting prodrugs that contain D-amino acid substituents; carbohydrate-cleaving enzymes such as beta-

galactosidase and neuraminidase useful for converting glycosylated prodrugs into free drugs; beta-lactamase useful for converting drugs derivatized with beta-lactams into free drugs; and penicillin amidases, such as penicillin V amidase or penicillin G amidase, useful for  
5 converting drugs derivatized at their amine nitrogens with phenoxyacetyl or phenylacetyl groups, respectively, into free drugs. Alternatively, antibodies with enzymatic activity, also known in the art as "abzymes", can be used to convert the prodrugs of the invention into free active drugs (see, e.g., Massey, *Nature* 328: 457-458 (1987)). Antibody-abzyme  
10 conjugates can be prepared as described herein for delivery of the abzyme to a tumor cell population.

The enzymes can be covalently bound to the antibodies by techniques well known in the art such as the use of heterobifunctional crosslinking reagents. Alternatively, fusion proteins comprising at least the antigen  
15 binding region of an antibody of the invention linked to at least a functionally active portion of an enzyme of the invention can be constructed using recombinant DNA techniques well known in the art (see, e.g., Neuberger et al., *Nature*, 312: 604-608 (1984)).

Further antibody modifications are contemplated. For example, the  
20 antibody may be linked to one of a variety of nonproteinaceous polymers, e.g., polyethylene glycol, polypropylene glycol, polyoxyalkylenes, or copolymers of polyethylene glycol and polypropylene glycol. The antibody also may be entrapped in microcapsules prepared, for example, by coacervation techniques or by interfacial polymerization (for example,  
25 hydroxymethylcellulose or gelatin-microcapsules and poly-(methylmethacrylate) microcapsules, respectively), in colloidal drug delivery systems (for example, liposomes, albumin microspheres, microemulsions, nano-particles and nanocapsules), or in macroemulsions. Such techniques are disclosed in *Remington's Pharmaceutical Sciences*,  
30 16th edition, Oslo, A., Ed., (1980). To increase the serum half life of the antibody, one may incorporate a salvage receptor binding epitope into the antibody (especially an antibody fragment) as described in U.S. Patent 5,739,277, for example. As used herein, the term "salvage receptor binding epitope" refers to an epitope of the Fc region of an IgG  
35 molecule (e.g., IgG<sub>1</sub>, IgG<sub>2</sub>, IgG<sub>3</sub>, or IgG<sub>4</sub>) that is responsible for increasing the *in vivo* serum half-life of the IgG molecule.

#### 7. Recombinant Methods

The invention also provides isolated nucleic acids encoding the antibodies as disclosed herein, vectors and host cells comprising the  
40 nucleic acid, and recombinant techniques for the production of the antibody.

For recombinant production of the antibody, the nucleic acid encoding it is isolated and inserted into a replicable vector for further cloning (amplification of the DNA) or for expression. DNA encoding the  
45 antibody is readily isolated and sequenced using conventional procedures

(e.g., by using oligonucleotide probes that are capable of binding specifically to genes encoding the antibody). Many vectors are available. The vector components generally include, but are not limited to, one or more of the following: a signal sequence, an origin of replication, one  
5 or more marker genes, an enhancer element, a promoter, and a transcription termination sequence.

The methods herein include methods for the production of chimeric or recombinant anti-Apo-2L receptor antibodies which comprise the steps of providing a vector comprising a DNA sequence encoding an anti-Apo-2L  
10 receptor antibody light chain or heavy chain (or both a light chain and a heavy chain), transfecting or transforming a host cell with the vector, and culturing the host cell(s) under conditions sufficient to produce the recombinant anti-Apo-2L receptor antibody product.

(i) *Signal sequence component*

15 The anti-Apo-2L receptor antibody of this invention may be produced recombinantly not only directly, but also as a fusion polypeptide with a heterologous polypeptide, which is preferably a signal sequence or other polypeptide having a specific cleavage site at the N-terminus of the mature protein or polypeptide. The heterologous signal sequence selected  
20 preferably is one that is recognized and processed (i.e., cleaved by a signal peptidase) by the host cell. For prokaryotic host cells that do not recognize and process the native antibody signal sequence, the signal sequence is substituted by a prokaryotic signal sequence selected, for example, from the group of the alkaline phosphatase, penicillinase, lpp,  
25 or heat-stable enterotoxin II leaders. For yeast secretion the native signal sequence may be substituted by, e.g., the yeast invertase leader,  $\alpha$  factor leader (including *Saccharomyces* and *Kluyveromyces*  $\alpha$ -factor leaders), or acid phosphatase leader, the *C. albicans* glucoamylase leader, or the signal described in WO 90/13646. In mammalian cell  
30 expression, mammalian signal sequences as well as viral secretory leaders, for example, the herpes simplex gD signal, are available. The DNA for such precursor region is ligated in reading frame to DNA encoding the antibody.

(ii) *Origin of replication component*

35 Both expression and cloning vectors contain a nucleic acid sequence that enables the vector to replicate in one or more selected host cells. Generally, in cloning vectors this sequence is one that enables the vector to replicate independently of the host chromosomal DNA, and includes origins of replication or autonomously replicating sequences.  
40 Such sequences are well known for a variety of bacteria, yeast, and viruses. The origin of replication from the plasmid pBR322 is suitable for most Gram-negative bacteria, the 2 $\mu$  plasmid origin is suitable for yeast, and various viral origins (SV40, polyoma, adenovirus, VSV or BPV) are useful for cloning vectors in mammalian cells. Generally, the origin  
45 of replication component is not needed for mammalian expression vectors

(the SV40 origin may typically be used only because it contains the early promoter).

(iii) Selection gene component

Expression and cloning vectors may contain a selection gene, also  
5 termed a selectable marker. Typical selection genes encode proteins that  
(a) confer resistance to antibiotics or other toxins, e.g., ampicillin,  
neomycin, methotrexate, or tetracycline, (b) complement auxotrophic  
deficiencies, or (c) supply critical nutrients not available from complex  
media, e.g., the gene encoding D-alanine racemase for *Bacilli*.

10 One example of a selection scheme utilizes a drug to arrest growth  
of a host cell. Those cells that are successfully transformed with a  
heterologous gene produce a protein conferring drug resistance and thus  
survive the selection regimen. Examples of such dominant selection use  
the drugs neomycin, mycophenolic acid and hygromycin.

15 Another example of suitable selectable markers for mammalian cells  
are those that enable the identification of cells competent to take up  
the antibody nucleic acid, such as DHFR, thymidine kinase,  
metallothionein-I and -II, preferably primate metallothionein genes,  
adenosine deaminase, ornithine decarboxylase, etc.

20 For example, cells transformed with the DHFR selection gene are  
first identified by culturing all of the transformants in a culture  
medium that contains methotrexate (Mtx), a competitive antagonist of  
DHFR. An appropriate host cell when wild-type DHFR is employed is the  
Chinese hamster ovary (CHO) cell line deficient in DHFR activity.

25 Alternatively, host cells (particularly wild-type hosts that  
contain endogenous DHFR) transformed or co-transformed with DNA sequences  
encoding the anti-Apo-2L receptor antibody, wild-type DHFR protein, and  
another selectable marker such as aminoglycoside 3'-phosphotransferase  
(APH) can be selected by cell growth in medium containing a selection  
30 agent for the selectable marker such as an aminoglycosidic antibiotic,  
e.g., kanamycin, neomycin, or G418. See U.S. Patent No. 4,965,199.

A suitable selection gene for use in yeast is the *trp1* gene present  
in the yeast plasmid YRp7 (Stinchcomb et al., *Nature*, 282:39 (1979)).  
The *trp1* gene provides a selection marker for a mutant strain of yeast  
35 lacking the ability to grow in tryptophan, for example, ATCC No. 44076 or  
PEP4-1. Jones, *Genetics*, 85:12 (1977). The presence of the *trp1* lesion  
in the yeast host cell genome then provides an effective environment for  
detecting transformation by growth in the absence of tryptophan.  
Similarly, *Leu2*-deficient yeast strains (ATCC 20,622 or 38,626) are  
40 complemented by known plasmids bearing the *Leu2* gene.

In addition, vectors derived from the 1.6  $\mu$ m circular plasmid pKD1  
can be used for transformation of *Kluyveromyces* yeasts. Alternatively,  
an expression system for large-scale production of recombinant calf  
chymosin was reported for *K. lactis*. Van den Berg, *Bio/Technology*, 8:135  
45 (1990). Stable multi-copy expression vectors for secretion of mature

recombinant human serum albumin by industrial strains of *Kluyveromyces* have also been disclosed. Fleer et al., *Bio/Technology*, 9:968-975 (1991).

(iv) Promoter component

5 Expression and cloning vectors usually contain a promoter that is recognized by the host organism and is operably linked to the antibody nucleic acid. Promoters suitable for use with prokaryotic hosts include the *phoA* promoter,  $\beta$ -lactamase and lactose promoter systems, alkaline phosphatase, a tryptophan (*trp*) promoter system, and hybrid promoters  
10 such as the *tac* promoter. However, other known bacterial promoters are suitable. Promoters for use in bacterial systems also will contain a Shine-Dalgarno (S.D.) sequence operably linked to the DNA encoding the anti-Apo-2L receptor antibody.

Promoter sequences are known for eukaryotes. Virtually all  
15 eukaryotic genes have an AT-rich region located approximately 25 to 30 bases upstream from the site where transcription is initiated. Another sequence found 70 to 80 bases upstream from the start of transcription of many genes is a CNCAAT region where N may be any nucleotide. At the 3' end of most eukaryotic genes is an AATAAA sequence that may be the signal  
20 for addition of the poly A tail to the 3' end of the coding sequence. All of these sequences are suitably inserted into eukaryotic expression vectors.

Examples of suitable promoting sequences for use with yeast hosts include the promoters for 3-phosphoglycerate kinase or other glycolytic  
25 enzymes, such as enolase, glyceraldehyde-3-phosphate dehydrogenase, hexokinase, pyruvate decarboxylase, phosphofructokinase, glucose-6-phosphate isomerase, 3-phosphoglycerate mutase, pyruvate kinase, triosephosphate isomerase, phosphoglucose isomerase, and glucokinase.

Other yeast promoters, which are inducible promoters having the  
30 additional advantage of transcription controlled by growth conditions, are the promoter regions for alcohol dehydrogenase 2, isocytochrome C, acid phosphatase, degradative enzymes associated with nitrogen metabolism, metallothionein, glyceraldehyde-3-phosphate dehydrogenase, and enzymes responsible for maltose and galactose utilization. Suitable  
35 vectors and promoters for use in yeast expression are further described in EP 73,657. Yeast enhancers also are advantageously used with yeast promoters.

Anti-Apo-2L receptor antibody transcription from vectors in  
mammalian host cells is controlled, for example, by promoters obtained  
40 from the genomes of viruses such as polyoma virus, fowlpox virus, adenovirus (such as Adenovirus 2), bovine papilloma virus, avian sarcoma virus, cytomegalovirus, a retrovirus, hepatitis-B virus and most preferably Simian Virus 40 (SV40), from heterologous mammalian promoters, e.g., the actin promoter or an immunoglobulin promoter, from heat-shock



promoters, provided such promoters are compatible with the host cell systems.

The early and late promoters of the SV40 virus are conveniently obtained as an SV40 restriction fragment that also contains the SV40 viral origin of replication. The immediate early promoter of the human cytomegalovirus is conveniently obtained as a HindIII E restriction fragment. A system for expressing DNA in mammalian hosts using the bovine papilloma virus as a vector is disclosed in U.S. Patent No. 4,419,446. A modification of this system is described in U.S. Patent No. 4,601,978. See also Reyes et al., Nature 297:598-601 (1982) on expression of human  $\beta$ -interferon cDNA in mouse cells under the control of a thymidine kinase promoter from herpes simplex virus. Alternatively, the rous sarcoma virus long terminal repeat can be used as the promoter.

(v) *Enhancer element component*

Transcription of a DNA encoding the anti-Apo-2L receptor antibody of this invention by higher eukaryotes is often increased by inserting an enhancer sequence into the vector. Many enhancer sequences are now known from mammalian genes (globin, elastase, albumin,  $\alpha$ -fetoprotein, and insulin). Typically, however, one will use an enhancer from a eukaryotic cell virus. Examples include the SV40 enhancer on the late side of the replication origin (bp 100-270), the cytomegalovirus early promoter enhancer, the polyoma enhancer on the late side of the replication origin, and adenovirus enhancers. See also Yaniv, Nature 297:17-18 (1982) on enhancing elements for activation of eukaryotic promoters. The enhancer may be spliced into the vector at a position 5' or 3' to the antibody-encoding sequence, but is preferably located at a site 5' from the promoter.

(vi) *Transcription termination component*

Expression vectors used in eukaryotic host cells (yeast, fungi, insect, plant, animal, human, or nucleated cells from other multicellular organisms) will also contain sequences necessary for the termination of transcription and for stabilizing the mRNA. Such sequences are commonly available from the 5' and, occasionally 3', untranslated regions of eukaryotic or viral DNAs or cDNAs. These regions contain nucleotide segments transcribed as polyadenylated fragments in the untranslated portion of the mRNA encoding the multivalent antibody. One useful transcription termination component is the bovine growth hormone polyadenylation region. See WO94/11026 and the expression vector disclosed therein.

(vii) *Selection and transformation of host cells*

Suitable host cells for cloning or expressing the DNA in the vectors herein are the prokaryote, yeast, or higher eukaryote cells described above. Suitable prokaryotes for this purpose include eubacteria, such as Gram-negative or Gram-positive organisms, for example, Enterobacteriaceae such as *Escherichia*, e.g., *E. coli*,

*Enterobacter*, *Erwinia*, *Klebsiella*, *Proteus*, *Salmonella*, e.g., *Salmonella typhimurium*, *Serratia*, e.g., *Serratia marcescans*, and *Shigella*, as well as *Bacilli* such as *B. subtilis* and *B. licheniformis* (e.g., *B. licheniformis* 41P disclosed in DD 266,710 published 12 April 1989),  
5 *Pseudomonas* such as *P. aeruginosa*, and *Streptomyces*. One optional *E. coli* cloning host is *E. coli* 294 (ATCC 31,446), although other strains such as *E. coli* B, *E. coli* X1776 (ATCC 31,537), and *E. coli* W3110 (ATCC 27,325) are suitable. These examples are illustrative rather than limiting.

10 In addition to prokaryotes, eukaryotic microbes such as filamentous fungi or yeast are suitable cloning or expression hosts for Apo-2L receptor antibody-encoding vectors. *Saccharomyces cerevisiae*, or common baker's yeast, is the most commonly used among lower eukaryotic host microorganisms. However, a number of other genera, species, and strains  
15 are commonly available and useful herein, such as *Schizosaccharomyces pombe*; *Kluyveromyces* hosts such as, e.g., *K. lactis*, *K. fragilis* (ATCC 12,424), *K. bulgaricus* (ATCC 16,045), *K. wickerhamii* (ATCC 24,178), *K. waltii* (ATCC 56,500), *K. drosophilum* (ATCC 36,906), *K. thermotolerans*, and *K. marxianus*; *Yarrowia* (EP 402,226); *Pichia pastoris* (EP 183,070);  
20 *Candida*; *Trichoderma reesia* (EP 244,234); *Neurospora crassa*; *Schwanniomyces* such as *Schwanniomyces occidentalis*; and filamentous fungi such as, e.g., *Neurospora*, *Penicillium*, *Tolypocladium*, and *Aspergillus* hosts such as *A. nidulans* and *A. niger*.

Suitable host cells for the expression of glycosylated antibody are  
25 derived from multicellular organisms. Examples of invertebrate cells include plant and insect cells. Numerous baculoviral strains and variants and corresponding permissive insect host cells from hosts such as *Spodoptera frugiperda* (caterpillar), *Aedes aegypti* (mosquito), *Aedes albopictus* (mosquito), *Drosophila melanogaster* (fruitfly), and *Bombyx mori* have been identified. A variety of viral strains for transfection  
30 are publicly available, e.g., the L-1 variant of *Autographa californica* NPV and the Bm-5 strain of *Bombyx mori* NPV, and such viruses may be used as the virus herein according to the present invention, particularly for transfection of *Spodoptera frugiperda* cells.

35 Plant cell cultures of cotton, corn, potato, soybean, petunia, tomato, and tobacco can also be utilized as hosts.

However, interest has been greatest in vertebrate cells, and propagation of vertebrate cells in culture (tissue culture) has become a routine procedure. Examples of useful mammalian host cell lines are  
40 monkey kidney CV1 line transformed by SV40 (COS-7, ATCC CRL 1651); human embryonic kidney line (293 or 293 cells subcloned for growth in suspension culture, Graham et al., *J. Gen Virol.* 36:59 (1977)); baby hamster kidney cells (BHK, ATCC CCL 10); Chinese hamster ovary cells/-DHFR (CHO, Urlaub et al., *Proc. Natl. Acad. Sci. USA* 77:4216 (1980));

mouse sertoli cells (TM4, Mather, *Biol. Reprod.* 23:243-251 (1980)); monkey kidney cells (CV1 ATCC CCL 70); African green monkey kidney cells (VERO-76, ATCC CRL-1587); human cervical carcinoma cells (HELA, ATCC CCL 2); canine kidney cells (MDCK, ATCC CCL 34); buffalo rat liver cells (BRL 3A, ATCC CRL 1442); human lung cells (W138, ATCC CCL 75); human liver cells (Hep G2, HB 8065); mouse mammary tumor (MMT 060562, ATCC CCL51); TRI cells (Mather et al., *Annals N.Y. Acad. Sci.* 383:44-68 (1982)); MRC 5 cells; FS4 cells; a human hepatoma line (Hep G2); and myeloma or lymphoma cells (e.g. Y0, J558L, P3 and NS0 cells) (see US Patent No. 5,807,715).

Host cells are transformed with the above-described expression or cloning vectors for antibody production and cultured in conventional nutrient media modified as appropriate for inducing promoters, selecting transformants, or amplifying the genes encoding the desired sequences.

(viii) *Culturing the host cells*

The host cells used to produce the antibody of this invention may be cultured in a variety of media. Commercially available media such as Ham's F10 (Sigma), Minimal Essential Medium ((MEM), (Sigma), RPMI-1640 (Sigma), and Dulbecco's Modified Eagle's Medium ((DMEM), Sigma) are suitable for culturing the host cells. In addition, any of the media described in Ham et al., *Meth. Enz.* 58:44 (1979), Barnes et al., *Anal. Biochem.* 102:255 (1980), U.S. Pat. Nos. 4,767,704; 4,657,866; 4,927,762; 4,560,655; or 5,122,469; WO 90/03430; WO 87/00195; or U.S. Patent Re. 30,985 may be used as culture media for the host cells. Any of these media may be supplemented as necessary with hormones and/or other growth factors (such as insulin, transferrin, or epidermal growth factor), salts (such as sodium chloride, calcium, magnesium, and phosphate), buffers (such as HEPES), nucleotides (such as adenosine and thymidine), antibiotics (such as GENTAMYCIN<sup>TM</sup> drug), trace elements (defined as inorganic compounds usually present at final concentrations in the micromolar range), and glucose or an equivalent energy source. Any other necessary supplements may also be included at appropriate concentrations that would be known to those skilled in the art. The culture conditions, such as temperature, pH, and the like, are those previously used with the host cell selected for expression, and will be apparent to the ordinarily skilled artisan.

(ix) *Purification*

When using recombinant techniques, the antibody can be produced intracellularly, in the periplasmic space, or directly secreted into the medium. If the antibody is produced intracellularly, as a first step, the particulate debris, either host cells or lysed fragments, is removed, for example, by centrifugation or ultrafiltration. Carter et al., *Bio/Technology* 10:163-167 (1992) describe a procedure for isolating antibodies which are secreted to the periplasmic space of *E. coli*. Briefly, cell paste is thawed in the presence of sodium acetate (pH 3.5), EDTA, and phenylmethylsulfonylfluoride (PMSF) over about 30 minutes. Cell

debris can be removed by centrifugation. Where the antibody is secreted into the medium, supernatants from such expression systems are generally first concentrated using a commercially available protein concentration filter, for example, an Amicon or Millipore Pellicon ultrafiltration unit. A protease inhibitor such as PMSF may be included in any of the foregoing steps to inhibit proteolysis and antibiotics may be included to prevent the growth of adventitious contaminants.

The antibody composition prepared from the cells can be purified using, for example, hydroxylapatite chromatography, gel electrophoresis, dialysis, and affinity chromatography, with affinity chromatography being the preferred purification technique. The suitability of protein A as an affinity ligand depends on the species and isotype of any immunoglobulin Fc region that is present in the antibody. Protein A can be used to purify antibodies that are based on human  $\gamma 1$ ,  $\gamma 2$ , or  $\gamma 4$  heavy chains (Lindmark et al., *J. Immunol. Meth.* 62:1-13 (1983)). Protein G is recommended for all mouse isotypes and for human  $\gamma 3$  (Guss et al., *EMBO J.* 5:1567-1575 (1986)). The matrix to which the affinity ligand is attached is most often agarose, but other matrices are available. Mechanically stable matrices such as controlled pore glass or poly(styrenedivinyl)benzene allow for faster flow rates and shorter processing times than can be achieved with agarose. Where the antibody comprises a  $C_H3$  domain, the Bakerbond ABX<sup>TM</sup> resin (J. T. Baker, Phillipsburg, NJ) is useful for purification. Other techniques for protein purification such as fractionation on an ion-exchange column, ethanol precipitation, Reverse Phase HPLC, chromatography on silica, chromatography on heparin SEPHAROSE<sup>TM</sup> chromatography on an anion or cation exchange resin (such as a polyaspartic acid column), chromatofocusing, SDS-PAGE, and ammonium sulfate precipitation are also available depending on the antibody to be recovered.

#### 30 C. FORMULATIONS

The Apo-2 ligand or Apo-2L receptor agonist antibody and CPT-11 are preferably administered in a carrier. The molecules can be administered in a single carrier, or alternatively, can be included in separate carriers. Suitable carriers and their formulations are described in Remington's Pharmaceutical Sciences, 16th ed., 1980, Mack Publishing Co., edited by Oslo et al. Typically, an appropriate amount of a pharmaceutically-acceptable salt is used in the carrier to render the formulation isotonic. Examples of the carrier include saline, Ringer's solution and dextrose solution. The pH of the solution is preferably from about 5 to about 8, and more preferably from about 7.4 to, about 7.8. It will be apparent to those persons skilled in the art that certain carriers may be more preferable depending upon, for instance, the route of administration and concentration of agent being administered. The carrier may be in the form of a lyophilized formulation or aqueous solution.

Acceptable carriers, excipients, or stabilizers are preferably nontoxic to cells and/or recipients at the dosages and concentrations employed, and include buffers such as phosphate, citrate, and other organic acids; antioxidants including ascorbic acid and methionine; 5 preservatives (such as octadecyldimethylbenzyl ammonium chloride; hexamethonium chloride; benzalkonium chloride, benzethonium chloride; phenol, butyl or benzyl alcohol; alkyl parabens such as methyl or propyl paraben; catechol; resorcinol; cyclohexanol; 3-pentanol; and m-cresol); low molecular weight (less than about 10 residues) polypeptides; 10 proteins, such as serum albumin, gelatin, or immunoglobulins; hydrophilic polymers such as polyvinylpyrrolidone; amino acids such as glycine, glutamine, asparagine, histidine, arginine, or lysine; monosaccharides, disaccharides, and other carbohydrates including glucose, mannose, or dextrans; chelating agents such as EDTA; sugars such as sucrose, 15 mannitol, trehalose or sorbitol; salt-forming counter-ions such as sodium; and/or non-ionic surfactants such as TWEEN™, PLURONICS™ or polyethylene glycol (PEG).

The formulation may also contain more than one active compound as necessary for the particular indication being treated, preferably those 20 with complementary activities that do not adversely affect each other. Alternatively, or in addition, the composition may comprise a cytotoxic agent, cytokine or growth inhibitory agent. Such molecules are suitably present in combination in amounts that are effective for the purpose intended.

25 The Apo-2L or agonist antibody and CPT-11 may also be entrapped in microcapsules prepared, for example, by coacervation techniques or by interfacial polymerization, for example, hydroxymethylcellulose or gelatin-microcapsules and poly-(methylmethacrylate) microcapsules, respectively, in colloidal drug delivery systems (for example, liposomes, 30 albumin microspheres, microemulsions, nano-particles and nanocapsules) or in macroemulsions. Such techniques are disclosed in Remington's Pharmaceutical Sciences 16th edition, Oslo, A. Ed. (1980).

The formulations to be used for in vivo administration should be sterile. This is readily accomplished by filtration through sterile 35 filtration membranes.

Sustained-release preparations may be prepared. Suitable examples of sustained-release preparations include semipermeable matrices of solid hydrophobic polymers containing the antibody, which matrices are in the form of shaped articles, e.g. films, or microcapsules. Examples of 40 sustained-release matrices include polyesters, hydrogels (for example, poly(2-hydroxyethyl-methacrylate), or poly(vinylalcohol)), polylactides (U.S. Pat. No. 3,773,919), copolymers of L-glutamic acid and  $\gamma$  ethyl-L-glutamate, non-degradable ethylene-vinyl acetate, degradable lactic acid-glycolic acid copolymers such as the LUPRON DEPOT™ (injectable 45 microspheres composed of lactic acid-glycolic acid copolymer and

leuprolide acetate), and poly-D-(-)-3-hydroxybutyric acid. While polymers such as ethylene-vinyl acetate and lactic acid-glycolic acid enable release of molecules for over 100 days, certain hydrogels release proteins for shorter time periods.

5 D. MODES OF ADMINISTRATION

The Apo-2L or Apo-2L receptor agonist antibody and CPT-11 can be administered in accord with known methods, such as intravenous administration as a bolus or by continuous infusion over a period of time, by intramuscular, intraperitoneal, intracerebrospinal, subcutaneous, intra-articular, intrasynovial, intrathecal, oral, topical, or inhalation routes. Optionally, administration may be performed through mini-pump infusion using various commercially available devices.

Effective dosages for administering Apo-2 ligand or agonist antibody and CPT-11 may be determined empirically, and making such determinations is within the skill in the art. It is presently believed that an effective dosage or amount of Apo-2 ligand used alone may range from about 1 µg/kg to about 100 mg/kg of body weight or more per day. An effective dosage or amount of CPT-11 used alone may range from about 1 mg/m<sup>2</sup> to about 150 mg/m<sup>2</sup>. Interspecies scaling of dosages can be performed in a manner known in the art, e.g., as disclosed in Mordenti et al., Pharmaceut. Res., 8:1351 (1991). Those skilled in the art will understand that the dosage of Apo-2 ligand or agonist antibody and CPT-11 that must be administered will vary depending on, for example, the mammal which will receive the Apo-2 ligand or agonist antibody and CPT-11, the route of administration, and other drugs or therapies being administered to the mammal.

Depending on the type of cells and/or severity of the disease, about 1 µg/kg to 15 mg/kg (e.g. 0.1-20 mg/kg) of agonist antibody is an initial candidate dosage for administration, whether, for example, by one or more separate administrations, or by continuous infusion. A typical daily dosage might range from about 1 µg/kg to 100 mg/kg or more, depending on the factors mentioned above. For repeated administrations over several days or longer, depending on the condition, the treatment is sustained until a desired suppression of disease symptoms occurs. However, other dosage regimens may be useful.

It is believed that pre-treatment of the cells with CPT-11 may reduce the amount of Apo-2L receptor agonist required to induce apoptosis in a selected population of cells. For example, pre-treatment of the cells with CPT-11 may reduce the amount of Apo-2L receptor agonist required to induce (an equivalent amount or degree of) apoptosis in the mammalian cells by at least 25% and preferably, by at least 50%.

It is contemplated that one or more Apo-2L receptor agonists may be employed in the methods. For example, the skilled practitioner may employ Apo-2 ligand, DR4 agonist antibody, DR5 agonist antibody, or

combinations thereof. Optionally, the Apo-2L receptor agonist antibody will comprise a cross-reactive antibody which binds to both DR4 and DR5.

It is contemplated that yet additional therapies may be employed in the methods. The one or more other therapies may include but are not  
5 limited to, other chemotherapies (or chemotherapeutic agents) and/or radiation therapy, immunoadjuvants, growth inhibitory agents, cytokines, and other non-Her-2 antibody-based therapies. Examples include interleukins (e.g., IL-1, IL-2, IL-3, IL-6), leukemia inhibitory factor, interferons, TGF-beta, erythropoietin, thrombopoietin, and anti-VEGF  
10 antibody. Other agents known to induce apoptosis in mammalian cells may also be employed, and such agents include TNF- $\alpha$ , TNF- $\beta$  (lymphotoxin- $\alpha$ ), CD30 ligand, 4-1BB ligand, and Apo-1 ligand.

Additional chemotherapies contemplated by the invention include chemical substances or drugs which are known in the art and are  
15 commercially available, such as Adriamycin, Doxorubicin, 5-Fluorouracil, Cytosine arabinoside ("Ara-C"), Cyclophosphamide, Leucovorin, Thiotepa, Busulfan, Cytosin, Taxol, Toxotere, Methotrexate, Cisplatin, Melphalan, Vinblastine, Bleomycin, Etoposide, Ifosfamide, Mitomycin C, Mitoxantrone, Vincristine, Vinorelbine, Carboplatin, Teniposide, Daunomycin,  
20 Carminomycin, Aminopterin, Dactinomycin, Mitomycins, Esperamicins (see U.S. Pat. No. 4,675,187), Melphalan and other related nitrogen mustards. Also included are agents that act to regulate or inhibit hormone action on tumors such as tamoxifen and onapristone.

Preparation and dosing schedules for such chemotherapy may be used  
25 according to manufacturers' instructions or as determined empirically by the skilled practitioner. Preparation and dosing schedules for such chemotherapy are also described in Chemotherapy Service Ed., M.C. Perry, Williams & Wilkins, Baltimore, MD (1992). The chemotherapeutic agent may precede, or follow administration with the Apo-2L or agonist antibody  
30 and/or CPT-11 or may be given simultaneously therewith.

The chemotherapy is preferably administered in a carrier, such as those described above. The mode of administration of the chemotherapy may be the same as employed for the Apo-2 ligand or agonist antibody or CPT-11 or it may be administered via a different mode.

35 Radiation therapy can be administered according to protocols commonly employed in the art and known to the skilled artisan. Such therapy may include cesium, iridium, iodine, or cobalt radiation. The radiation therapy may be whole body irradiation, or may be directed locally to a specific site or tissue in or on the body. Typically,  
40 radiation therapy is administered in pulses over a period of time from about 1 to about 2 weeks. The radiation therapy may, however, be administered over longer periods of time. Optionally, the radiation therapy may be administered as a single dose or as multiple, sequential doses.

Following administration of Apo-2 ligand or agonist antibody and CPT-11, treated cells *in vitro* can be analyzed. Where there has been *in vivo* treatment, a treated mammal can be monitored in various ways well known to the skilled practitioner. For instance, tumor mass may be  
5 observed physically, by biopsy or by standard x-ray imaging techniques.

### III. Articles of Manufacture

In another embodiment of the invention, an article of manufacture containing materials useful for the treatment of the disorders described above is provided. The article of manufacture comprises a container and a label. Suitable containers include, for example, bottles, vials, syringes, and test tubes. The containers may be formed from a variety of materials such as glass or plastic. The container holds a composition which is effective for treating the condition and may have a sterile  
15 access port (for example the container may be an intravenous solution bag or a vial having a stopper pierceable by a hypodermic injection needle). The active agents in the composition are the Apo-2 ligand or agonist antibody and CPT-11. The label on, or associated with, the container indicates that the composition is used for treating the condition of choice. The article of manufacture may further comprise a second  
20 container comprising a pharmaceutically-acceptable buffer, such as phosphate-buffered saline, Ringer's solution and dextrose solution. It may further include other materials desirable from a commercial and user standpoint, including other buffers, diluents, filters, needles, syringes, and package inserts with instructions for use.  
25

The following examples are offered by way of illustration and not by way of limitation. The disclosures of all citations in the specification are expressly incorporated herein by reference.

30

#### EXAMPLE 1

This example illustrates the synergistic inhibition of tumor growth by Apo-2 ligand and CPT-11 *in vivo*.

The colon carcinoma cell line COLO205 (available from NCI) were grown and maintained according to the supplier's methods. Briefly,  
35 COLO205 cells were cultured in high glucose DMEM/F12 (50:50) media containing 10% fetal bovine serum and 2.0mM L-Glutamine. Apo-2 ligand comprising amino acids 114-281 (SEQ ID NO:1) was prepared in *E. coli*. The extracellular portion of human Apo-2L (amino acids 114-281; see Pitti et al., supra) was subcloned into the pS1346 expression plasmid  
40 (Scholtissek et al., Gene, 62:55-64 (1988)) with an added initiator methionine codon, and expressed under control of the *trp* promoter in *E. coli* strain W3110, in 10L or 100 L fermentors. Cell-paste containing recombinant human soluble Apo-2L was extracted with a buffer containing 0.1M Tris/0.2M NaCl/50 mM EDTA, pH 8.0. The extract was precipitated by  
45 40% ammonium sulfate. Purification to >98 % homogeneity was achieved by



two consecutive chromatographic separation steps on hydroxyapatite and Ni-NTA agarose columns. (Although it lacks a polyhistidine tag, the recombinant soluble 114-281 amino acid Apo-2L fragment is believed to bind to the Ni-NTA column through endogenous histidine residues). Purity was determined by sodium-dodecyl sulfate (SDS)-polyacrylamide gel electrophoresis and silver-nitrate or coomassie-blue staining, by amino acid sequence analysis, and by size-exclusion on high performance liquid chromatography (HPLC). CPT-11 (Camptosar®) was obtained from Pharmacia & Upjohn.

Athymic nude mice (Jackson Laboratories) were injected subcutaneously with 5 million COLO205 colon carcinoma cells and the tumors allowed to grow to about 120 mm<sup>3</sup>. Tumor-bearing mice were randomized into 4 groups at 9 mice per group and treated with either vehicle (20mM Tris, 8% Trehalose, 0.01 Tween-20, pH 7.5), Apo-2L (30 mg/kg/day on days 0-4 and 7-11), or CPT-11 (80 mg/kg/day on days 0, 4, and 8), or a combination of Apo-2L (30 mg/kg/day on days 0-4 and 7-11) plus CPT-11 (80 mg/kg/day on days 0, 4, and 8). Tumor volumes were determined at the indicated days over 34 days.

As shown in Figure 1, Apo-2L (open triangles) or CPT-11 (open squares) each suppressed tumor growth during the treatment period, although tumor growth resumed several days later in all 9 animals of each group. In contrast, the combination of Apo-2L with CPT-11 (closed triangles) caused substantial tumor shrinkage, resulting in complete tumor elimination in 8 out of 9 animals in the combination treatment group.

The results of this experiment indicate that combinations of Apo-2 ligand and CPT-11 treatment synergistically inhibited growth of cancer cells *in vivo*.

## EXAMPLE 2

This example illustrates the synergistic inhibition of tumor growth by the DR4 receptor agonist antibody, 4H6.17.8 ("4H6"), and CPT-11 *in vivo*.

The agonist antibody was prepared as follows. A soluble DR4 ECD immunoadhesin construct was prepared. A mature DR4 ECD sequence (amino acids 1-218 shown in Pan et al., *supra*) was cloned into a pCMV-1 Flag vector (Kodak) downstream of the Flag signal sequence and fused to the CH1, hinge and Fc region of human immunoglobulin G<sub>1</sub> heavy chain as described previously [Aruffo et al., *Cell*, 61:1303-1313 (1990)]. The immunoadhesin was expressed by transient transfection into human 293 cells and purified from cell supernatants by protein A affinity chromatography, as described by Ashkenazi et al., *Proc. Natl. Acad. Sci.*, 88:10535-10539 (1991).

Balb/c mice (obtained from Charles River Laboratories) were immunized by injecting 0.5 µg/50 µl of a DR4 ECD immunoadhesin protein (as

described above)(diluted in MPL-TDM adjuvant purchased from Ribi Immunochemical Research Inc., Hamilton, MT) 11 times into each hind foot pad at 3-4 day intervals.

Three days after the final boost, popliteal lymph nodes were removed from the mice and a single cell suspension was prepared in DMEM media (obtained from Biowhitakker Corp.) supplemented with 1% penicillin-streptomycin. The lymph node cells were then fused with murine myeloma cells P3X63AgU.1 (ATCC CRL 1597) using 35% polyethylene glycol and cultured in 96-well culture plates. Hybridomas resulting from the fusion were selected in HAT medium. Ten days after the fusion, hybridoma culture supernatants were screened in an ELISA to test for the presence of monoclonal antibodies binding to the DR4 ECD immunoadhesin protein (described above).

In the ELISA, 96-well microtiter plates (Maxisorp; Nunc, Kamstrup, Denmark) were coated by adding 50  $\mu$ l of 2  $\mu$ g/ml goat anti-human IgG Fc (purchased from Cappel Laboratories) in PBS to each well and incubating at 4°C overnight. The plates were then washed three times with wash buffer (PBS containing 0.05% Tween 20). The wells in the microtiter plates were then blocked with 200  $\mu$ l of 2.0% bovine serum albumin in PBS and incubated at room temperature for 1 hour. The plates were then washed again three times with wash buffer.

After the washing step, 50  $\mu$ l of 0.4  $\mu$ g/ml DR4 ECD immunoadhesin protein in assay buffer was added to each well. The plates were incubated for 1 hour at room temperature on a shaker apparatus, followed by washing three times with wash buffer.

Following the wash steps, 100  $\mu$ l of the hybridoma supernatants or Protein G-sepharose column purified antibody (10  $\mu$ g/ml) was added to designated wells. 100  $\mu$ l of P3X63AgU.1 myeloma cell conditioned medium was added to other designated wells as controls. The plates were incubated at room temperature for 1 hour on a shaker apparatus and then washed three times with wash buffer.

Next, 50  $\mu$ l HRP-conjugated goat anti-mouse IgG Fc (purchased from Cappel Laboratories), diluted 1:1000 in assay buffer (0.5% bovine serum albumin, 0.05% Tween-20 in PBS), was added to each well and the plates incubated for 1 hour at room temperature on a shaker apparatus. The plates were washed three times with wash buffer, followed by addition of 50  $\mu$ l of substrate (TMB Microwell Peroxidase Substrate; Kirkegaard & Perry, Gaithersburg, MD) to each well and incubation at room temperature for 10 minutes. The reaction was stopped by adding 50  $\mu$ l of TMB 1-Component Stop Solution (Diethyl Glycol; Kirkegaard & Perry) to each well, and absorbance at 450 nm was read in an automated microtiter plate reader.

Hybridoma supernatants initially screened in the ELISA were considered for their ability to bind to DR4-IgG but not to CD4-IgG. The

supernatants testing positive in the ELISA were further analyzed by FACS analysis using 9D cells (a human B lymphoid cell line expressing DR4; Genentech, Inc.) and FITC-conjugated goat anti-mouse IgG. For this analysis, 25  $\mu$ l of cells suspended (at  $4 \times 10^6$  cells/ml) in cell sorter buffer (PBS containing 1% FCS and 0.02%  $\text{NaN}_3$ ) were added to U-bottom microtiter wells, mixed with 100  $\mu$ l of culture supernatant or purified antibody (10  $\mu$ g/ml) in cell sorter buffer, and incubated for 30 minutes on ice. The cells were then washed and incubated with 100  $\mu$ l FITC-conjugated goat anti-mouse IgG for 30 minutes at 4°C. Cells were then washed twice, resuspended in 150  $\mu$ l of cell sorter buffer and then analyzed by FACScan (Becton Dickinson, Mountain View, CA).

The FACS staining of the 9D cells revealed that the antibodies, 4E7.24.3 and 4H6.17.8, recognized the DR4 receptor on the 9D cells. Hybridoma supernatants and purified antibodies were then tested for activity to induce DR4 mediated 9D cell apoptosis. The 9D cells ( $5 \times 10^5$  cells/0.5ml) were incubated with 5  $\mu$ g of DR4 mAbs (4E7.24.3 or 4H6.17.8) or IgG control antibodies in 200  $\mu$ l complete RPMI media at 4°C for 15 minutes. The cells were then incubated for 5 minutes at 37°C with or without 10  $\mu$ g of goat anti-mouse IgG Fc antibody (ICN Pharmaceuticals) in 300  $\mu$ l of complete RPMI. At this point, the cells were incubated overnight at 37°C and in the presence of 7%  $\text{CO}_2$ . The cells were then harvested and washed once with PBS. The apoptosis of the cells was determined by staining of FITC-annexin V binding to phosphatidylserine according to manufacturer recommendations (Clontech). The cells were washed in PBS and resuspended in 200  $\mu$ l binding buffer. Ten  $\mu$ l of annexin-V-FITC (1  $\mu$ g/ml) and 10  $\mu$ l of propidium iodide were added to the cells. After incubation for 15 minutes in the dark, the 9D cells were analyzed by FACS.

Both DR4 antibodies (in the absence of the goat anti-mouse IgG Fc) induced apoptosis in the 9D cells as compared to the control antibodies. Agonistic activity of both DR4 antibodies, however, was enhanced by DR4 receptor cross-linking in the presence of the goat anti-mouse IgG Fc. This enhanced apoptosis by both DR4 antibodies is comparable to the apoptotic activity of Apo-2L in 9D cells.

The *in vivo* study examining the effects of the 4H6.17.8 monoclonal antibody plus CPT-11 (as compared to other treatment groups indicated in Figure 2) was conducted essentially as described in Example 1 above, except that in the antibody treatment groups, anti-DR4 antibody 4H6 (5 mg/kg; prepared as described above) was administered by *i.p.* injection to the mice twice per week for the duration of the study. In the Apo-2L treatment groups, Apo-2L was administered by *i.p.* injection on days 0-4

at 60 mg/kg/day. In the CPT-11 treatment groups, CPT-11 was administered by i.v. injection on days 0, 4, and 8 at 80 mg/kg.

The results are shown in Figure 2. Each agent alone caused a significant delay in tumor progression. The combination of Apo-2L or anti-DR4 monoclonal antibody with CPT-11 caused tumor regression, with a much more delayed time to tumor progression as compared to the single agent treatments. The anti-DR4 monoclonal antibody was more effective than Apo-2L both as single agent and in combination with CPT-11. A partial response (tumor volume decreased by more than 50% of its initial value) occurred in all 10 mice treated with the anti-DR4 antibody plus CPT-11, but in only 6 out of 10 mice treated with the Apo-2L plus CPT-11. These results show that Apo-2L receptor agonists cooperate synergistically with CPT-11 to inhibit tumor progression beyond the additive sum of effects of the respective single agent treatments.

### EXAMPLE 3

This example describes physiological effects of CPT-11 and Apo2L/TRAIL and demonstrates how pre-treatment of cells with CPT-11 prior to exposure to Apo2L/TRAIL produces the highest induction of DR5 and DR4 mRNA, as well as caspase -3- like cleavage/activation and apoptosis.

The abbreviations used herein include: Apo2L/TRAIL, Apo2 ligand/tumor necrosis factor related apoptosis-inducing ligand (prepared as described in Examples 1 and 2); DR, death receptor; DcR, decoy receptor; FADD, Fas-associating protein with death domain; CPT, Camptothecin; CPT-11, irinotecan; HUVEC, human umbilical vein endothelial cells; TNF, tumor necrosis factor; FLIP, fllice-inhibitory protein; CDDP, cis-diamminedichloroplatinum (II); CDK, cyclin-dependent kinase.

For cell culture, the human tumor colon cancer cell line HCT116 was obtained from the American Type Culture Collection (Manassas, VA). Cells were cultured in RPMI 1640 medium with 10% fetal bovine serum, 1 mM Glutamine, 100 units/ml penicillin and 100 µg/ml streptomycin. Cells were subcultured in 150 cm plates 24 hours before drug treatment. Human umbilical vein endothelial cells (HUVEC) were obtained from Cell Systems (#2V0-C75; Kirkland, WA) and incubated in CS-C™ complete medium (#4Z0-500, Cell Systems).

An AlamarBlue™ assay was used to determine the cell viability. HCT116 colon cancer cells (~10000 cells/well) were incubated overnight in 10% FBS RPMI 1640 medium in 96-well tissue culture plates. The medium was removed the following day, and the cells were incubated for 24 hours in serum free medium with Apo2L/TRAIL alone (1 µg/mL), CPT-11 (50 µg/mL), or Apo2L/TRAIL + CPT 11. AlamarBlue™ was added to the wells for the last 6 hours of the 24 hours incubation time. Fluorescence was read using 96-well fluorometer plate reader with an excitation of 530 nm and emission of 590 nm (CytoFluor multi-well plate reader series 4000, PerSeptive Biosystems; Framingham, MA). In addition, the Molecular

probes Live/Dead<sup>R</sup> viability/cytotoxicity kit (Eugene, OR) was used to evaluate the presence of live or dead cells. In these assays, Calcein-Am (4  $\mu$ M) and ethidium homodimer-1 (2  $\mu$ M) were added to treated cells 15 minutes before inspecting the cultures under an Axiovert 25 (Zeiss; Thornwood, NY) fluorescence microscope equipped with fluorescein (calcein) and rhodamine (ethidium homodimer-1) filters.

A crystal violet assay was also used. HCT116 colon cancer cells (approximately 20,000 cells/well) were incubated overnight in 10% fetal bovine serum (FBS) RPMI 1640 medium in 96-well tissue culture plates. The medium was removed the following day, and the cells were incubated for 24 or 48 hours in the fresh medium containing the various agents noted above. At the end of each treatment, the medium was removed and 100  $\mu$ l of 0.5% crystal violet solution was added to each well and incubated at room temperature for 10 minutes before washing with water. After the wells were dry, 100  $\mu$ l of ethanol containing 0.5 N HCL was added to each well. The plates were then read at 540 nm using a 96-well plate reader (Spectra Max 340pc, Molecular Devices Corporation, Sunnyvale, CA).

Caspase activity was determined by caspase-3 assay kits (Clontech; Palo Alto, CA). The assay was performed according to manufacturer's instruction. HCT116 cells were cultured in RPMI medium containing 10% FCS, 1 mM Glutamine, 100 units/ml penicillin, and 100  $\mu$ g/ml streptomycin and subjected to various treatments as described in the figure legends. After treatment, cells were collected, washed with cold PBS once, and frozen at -20°C until the time of assay. The cell pellets were thawed and lysed on ice for 10 minutes by the cell lysis buffer provided in the kit. The lysates were incubated with the fluorogenic caspase substrate (Z-DEVD-AFE, 100  $\mu$ M) in reaction buffer at 37°C for one hour. The samples were analyzed in a CytoFluor multi-well plate reader (PerSeptive Biosystems) with a 400/30 nm excitation filter and a 508/20 nm emission filter. The levels of relative fluorescence were normalized against the protein concentration of each sample.

Total RNA was isolated using the RNA Stat-60<sup>TM</sup> solution (Tel-Test, Inc (Friendsweed, TX) according to manufacturer's instruction. The Quantigene bDNA<sup>TM</sup> signal amplification kit (Chiron diagnostics; East Walpole, MA) was used to evaluate mRNA levels. The sequence of GAPDH probe sets was as recommended by the manufacturer. The DR4 probe sets were lined within the region of nucleic acid residues 9 to 582. There were 5 capture probes, 16 labeling probes and 8 blocking probes. The DR5 probe sets were lined within the region of nucleic acid residues 13 - 591. There were 5 capture probes, 17 labeling probes and 5 blocking probes for DR5. The specificity of the probes was tested using RNA from in vitro transcription using recombinant DcR1, DcR2, DR4 and DR5 constructs. Both DR4 and DR5 probe sets were highly specific for their

own RNA transcripts. The signal from each probe sets was linear with the concentrations tested. bDNA assays were performed according to manufacturer's instructions using about 2 µg of total RNA/well.

For Western blotting, cells were lysed in 20 mM Tris-HCl, pH 7.4  
5 containing 10% glycerol, 1 % Triton-x100, 150 mM NaCl and protease inhibitors. (1 mM PMSF, 10 µg/ml Aprotinin, 10 µg/ml Leupeptin, Sigma). Aliquots of 50 µg of total protein per well were separated in a NuPAGE 4-12% Bis-Tris gel with NuPAGE MES SDS running buffer (Novex; San Diego, CA). The gels were transferred to 0.2 µm pure nitrocellulose membrane  
10 (Bio-Rad) by semi-dry transfer cell (Bio-dad; Hercules, CA) with NuPAGE transfer buffer. The membranes were blocked with PBS containing 5% nonfat dry milk and incubated with primary antibodies against the proteins of interest followed by a horseradish peroxide-coupled secondary antibody (Amersham; Braunschweig, Germany). Immunoreactive bands were  
15 visualized by the enhanced chemiluminescence system (Amersham; Braunschweig, Germany). The antibodies used in this study were: Anti-caspase-3 (Stratagene, #200021; La Jolla, CA), anti-p53 and anti-p21 (Oncogene, #OP43 and #OP64; Cambridge, MA), and anti-FLIP (#343002, Calbiochem, San Diego, CA). The antibodies were used at the  
20 concentrations recommended by the manufacturers.

Cell cycle analysis was performed via FACS analysis. HCT116 and HUVEC cells were incubated and treated as described in the figure legends. After the treatments, both floating cells in the medium and  
live cells in the plate were collected. Cycle TEST PLUS DNA™ reagent  
25 kit (Beckon Dickinson; San Jose, CA) was used to stain cells according to manufacturer's instruction. The stained cells were analyzed in a FACS sorter (Becton Dickinson). The percentage of apoptotic cells containing a sub-G1 DNA content was quantitated using the CellQuest program. The  
percentage of live cells in each phase of cell cycle was quantitated  
30 using the ModFit LT program.

Figure 5 provides an apoptotic time profile showing that the CPT-11 sensitization of Apo2L/TRAIL-mediated apoptosis of HCT116 cells *in vitro* was time dependent. The tumor cells were stained with calcein-AM and ethidium homodimer-1 (green and red fluorescence, respectively) following  
incubation with Apo2L/TRAIL or CPT-11 alone, and in combinations. Under  
35 these conditions, green fluorescence depicts living cells; and red staining is indicative of dead cells. After 2 hours of treatment, Apo2L/TRAIL and the Apo2L/TRAIL + CPT-11 combination induced cellular changes characteristic of apoptosis, including cell shrinkage and  
40 cellular detachment from the monolayer. However, there were no noticeable differences in cell killing between Apo2L/TRAIL and Apo2L/TRAIL + CPT-11 at this time. In contrast, the combination of Apo2L/TRAIL + CPT-11, resulted in a clear increase in cell death by 24  
hours, as demonstrated by the uptake of ethidium homodimer and by the  
45 evident decrease in total cell density. Incubation with CPT-11 alone did

not show any morphological changes at 2 hours. However, a few dead cells were clearly present after 24 hours of treatment. Results from a quantitative analysis of cell survival (Figure 4) are consistent with the morphological fluorescent data. At 24 hours of treatment, the combination of Apo2L/TRAIL + CPT-11 resulted in a 26 % increase in apoptosis in comparison with Apo2L alone.

In the crystal violet assay conducted, some HCT116 cells were exposed to the combination treatment Apo2L/TRAIL (100ng/m) and CPT-11 (50 microgram/ml) for a total of 24 hours, followed for another 24 hour incubation in the presence of medium alone. In the group of cells treated sequentially, the HCT116 cells were exposed for the initial 24 hours to CPT-11, then the medium having been changed, were exposed to Apo2L/TRAIL containing medium for another 24 hours. In these conditions, the total cell killing in the cells treated sequentially was enhanced by about 6% above the cell samples treated with the combination treatment ( $p < 0.001$ , t-Test). Moreover, the relative survival activity comparing the sequential and combination treatments decreased by as much as 54%. This effect was observed at different concentrations of CPT-11 and Apo2L/TRAIL (data not shown).

As it has been previously reported that Apo2L/TRAIL-induced apoptosis involves caspase-3 activity (see e.g. Muhlenbeck et al., J.B.C. 273: 33091-8 (1998)), levels of caspase-3 activation were measured under these conditions. Specifically, the assessment of caspase-3 activity over time was monitored by fluorometric and western blot analysis as described above. Western blot analysis of caspase-3 activation showed that after 2 hours of treatment, Apo2L/TRAIL induced significant cleavage of caspase-3 into its p24, p20, and p17 forms (Figure 5). Caspase-3 activation was confirmed independently by a fluorometric assay (Figure 5). Interestingly, the combination of Apo2L/TRAIL + CPT-11 induced a similar degree of caspase-3 cleavage and activity after 2 hours treatment. CPT-11 alone induced a small but noticeable caspase-3-like activity, but cleavage was undetectable on Western blots. At 24 hours, the combined incubation of Apo2L/TRAIL + CPT-11 caused a clear increase in caspase-3 processing and activity in comparison with Apo2L/TRAIL or CPT-11 alone. Furthermore, a variation of the combination treatment in which the cells were incubated overnight with CPT-11 alone, followed by 2 hours of treatment with Apo2L/TRAIL in addition to CPT-11, resulted in the highest degree of caspase-3 cleavage and activity. Specifically, the pretreatment of cells with CPT-11 for 20-22 hours followed by two hours with Apo2L/TRAIL produced the highest induction of DR5 and DR4 mRNA, as well as caspase -3- like cleavage/activation and apoptosis. Taken together, these results show an enhancement in caspase-3 activation after combined Apo2L/TRAIL + CPT-11 treatment that leads to increased tumor apoptosis.

To investigate the effects of Apo2L/TRAIL and CPT-11 on the expression of Apo2L/TRAIL receptors DR5 and DR4, a bDNA assay was used. HCT116 cells were analyzed before and after treatment with Apo2L/TRAIL or CPT-11 alone, and in combination. Apo2L/TRAIL induced a two-fold transient increase in DR5 mRNA expression compared with controls after 2 hours of treatment (Figure 6). Apo2L/TRAIL-induced changes in DR5 expression returned to control levels after 24 hours of treatment. In contrast, CPT-11 alone resulted in a 2.5-fold increase in both DR4 and DR5 mRNA after 24 hours of treatment but not at 2 hours (Figure 6). Treatment with CPT-11 alone for 22 hours, followed by treatment with Apo2L/TRAIL + CPT-11 for another 2 hours resulted in the highest upregulation of DR5 expression (3 to 4-fold). All of the treatments for 30 minutes resulted in no changes in receptor expression. The time profile of the levels of caspase-3 cleavage/activation followed the upregulation of DR5 and/or DR4 at 2 and 24 hours, respectively (Figure 5). In contrast, DR5 and DR4 expression in HUVEC cells was not affected by any of these treatments. These results suggest that the upregulation of DR5 in tumor cells by Apo2L/TRAIL may serve to enhance its own apoptotic activity. The further upregulation of both DR5 and DR4 by the combined Apo2L/TRAIL + CPT-11 treatment supports this observation. In addition, a similar set of experiments was performed in the presence of a general caspase inhibitor, Z-VAD, to analyze the entire cell population rather than the surviving cells at 24 hours. The combination Apo2L/TRAIL + CPT-11 with Z-VAD resulted in an additional increase (1.7 versus 2.7 fold for DR5 and 1.9 versus 3.0 fold increase for DR4) compared to respective controls without Z-VAD (Figure 7). These results are in agreement with the idea that upregulation of these death receptors contributes to enhanced cell death.

To determine the involvement of p53 in DR5 upregulation by Apo2L/TRAIL and CPT-11, p53 protein expression levels were measured by western blot analysis. Aliquots were analyzed after HCT116 tumor and normal HUVEC cells were treated with Apo2L/TRAIL or CPT-11 alone, and in combination. Consistent with previous reports indicating that Apo2L/TRAIL-mediated apoptosis is p53 independent (see e.g. Ashkenazi et al., Current Opinion in Cell Biology 11: 255-260 (1999) and Rieger et al., FEBS Letters 427: 124-128 (1998)) Apo2L/TRAIL did not increase p53 protein level at any time-point analyzed (Figure 8). In contrast, as previously reported (McDonald et al., British Journal of Cancer 78:745-51 (1998)), incubation with CPT-11 resulted in a strong and sustained induction of p53 expression as early as 2 hours of treatment in both tumor and normal cells. CPT-11 induction of p53 persisted for at least 24 hours, and this induction was not affected by the addition of Apo2L/TRAIL.

The role of p21 in apoptosis versus cell arrest was also examined. Aliquots were analyzed in parallel for p21 and p53 protein levels by



western blot. As previously shown, CPT-11 alone strongly induced p53 in both cell types (Figure 8). CPT-11 also mediated a large induction of p21 protein at 24 hours both in tumor and normal cells (Figure 9). Apo2L/TRAIL alone did not have an effect on p53 or p21 expression. The combination treatment of Apo2L/TRAIL + CPT-11 also induced strong p53 and p21 expression after 24 hours of treatment in normal cells. Surprisingly, the levels of p21 protein in the combination treatment of HCT116 tumor cells remained at baseline levels regardless of the increase in p53 expression similar in magnitude to the CPT-11 alone. These data provided evidence that Apo2L/TRAIL suppresses the accumulation of p21 associated with the increase in p53 after CPT-11 treatment.

The possible involvement of FLIP in the experimental model of tumor apoptosis *in vitro* was also investigated. HCT116 cells were treated as previously described for 2 and 24 hours. Cell lysates were obtained and processed for western blot analysis using anti-FLIP antibodies. Figure 10 shows that the protein levels of FLIP were unaffected by any experimental treatment. (This indicates that FLIP was not a factor in the regulation of apoptosis using this colon carcinoma cell line).

To determine a direct correlation between p21 induction and changes in the cell cycle profile of treated cells and in particular, the appearance of cell cycle arrest, HCT116 cells were subjected to cell cycle analysis after 2, 6, and 24 hours of treatment with Apo2L/TRAIL or CPT-11 alone, and in combination. At 6 hours (Figure 11), CPT-11 alone induced a significant shift in the cell cycle profile resulting in a Go-G1 cell cycle arrest (76%). This change was also present, albeit to a lesser degree (55%) in the combination Apo2L/TRAIL + CPT-11 treatment and was not induced by Apo2L/TRAIL alone (43% vs. 30% control). By 24 hours, CPT-11 treatment resulted in an entirely different profile characterized by the appearance of a G2-M phase arrest (43% vs. 19% control and a clear reduction in Go-G1 phase. More importantly, the combination of Apo2L/TRAIL + CPT-11 completely prevented the appearance of this G2-M arrest (17% vs. 19% control) after 24 hours of treatment. These data are consistent with the Apo2L/TRAIL-mediated suppression of CPT-11 induction of p21.

Cell cycle analysis of normal cells for 24 hours under similar experimental conditions showed no differences among the treatments. These cell cycle analyses also provided confirmation of the increased apoptotic activity of the combination Apo2L/TRAIL + CPT-11 treatment. Apo2L/TRAIL + CPT-11 (24 hours) resulted in 42 % apoptosis, 19% with Apo2L/TRAIL alone, and 9% with CPT-11 treatment in comparison to 1% in control cells (Figure 11). These results further support the concept that the combined Apo2L/TRAIL + CPT-11 treatment mediates tumor suppression by preventing p21 induction and directing the cancer cells towards the apoptotic, rather than the cell cycle arrest pathway. A more detailed time profile of changes in p21 protein levels by CPT-11

incubation indicated increases as early as 4 hours of treatment that increased further by 18 hours of treatment.

Recent studies have indicated opposite roles for p21 in the apoptotic process, either as a caspase substrate (see e.g. Zhang et al., Oncogene 18:1131-1138 (1999); Levkau et al., Molecular Cell 1: 553-563 (1998); Gervais et al., Journal of Biological Chemistry 273: 19207-19212 (1998)) or as an inhibitor of caspase activation (see e.g. Suzuki et al., Oncogene 17: 931-939 (1998); Suzuki et al., Oncogene 18: 1239-44 (1999); Suzuki et al., Molecular & Cellular Biology 19:3842-3847 (1999); Suzuki et al., Oncogene 19: 1346-1353 (2000)). To determine the role of caspase Apo2L/TRAIL in the protein levels of p21, HCT116 cells were treated as previously described but also in the presence of the general caspase inhibitor, Z-VAD. Caspase inhibition resulted in a noticeable increase in the cellular levels of p21 in the combination Apo2L/TRAIL + CPT-11 group as compared to the same treatment without Z-VAD (Figure 12). Major differences were not observed with the remaining treatment groups. Interestingly, when tumor cells were pre-incubated with CPT-11 overnight and then treated for 2 hours with Apo2L/TRAIL before analysis, there was a similar decrease in p21 levels as detected in the regular combination treatment for 24 hours. Degradation of p21 was confirmed by the presence of a cleaved fragment at approximately 15 Kd under these conditions. This experiment demonstrated that inhibition of caspase activity prevented the otherwise strong degradation of p21 induced by Apo2L/TRAIL. To further show a functional correlation between caspase activation, levels of p21 and cell cycle arrest, cell cycle analysis was performed and showed that Apo2L/TRAIL did not prevent the CPT-11-induced G2-M arrest in the presence of the caspase inhibitor Z-VAD (Figure 13).

Further experiments were conducted to examine combination regimens for Apo2L/TRAIL and CPT-11, and two different conditions of the combination treatment were compared. In the first condition (combination), tumor cells were exposed to Apo2L/TRAIL and CPT-11 in vitro for a total of 24 hours, followed for another 24 hours incubation in the presence of medium alone. In the second group (sequential), cells were exposed for the initial 24 hours to CPT-11, then changed to Apo2L/TRAIL alone containing medium for another 24 hours. Figure 14A shows that the total cell killing in the sequential treatment was enhanced about 6% more than the combination group ( $p < 0.001$ , t-Test). Moreover, the relative killing activity comparing the sequential and combination treatments increased as much as 54%. This effect was seen at different concentrations of Apo2L/TRAIL (Figure 14A). Furthermore, this increased apoptotic effect of the sequential over the combination treatment was further enhanced as much as 68% when the cells were incubated for additional four days in drug free medium following 2 days of drug exposure (Figure 14B).

Increased tumor cell death was also observed in the combination and sequential treatments when the active metabolite of CPT-11, SN38, was used instead (Figure 15), indicating that the increase in tumor apoptosis does not reflect changes in the metabolism of CPT-11 compound.

5

#### Deposit of Material

The following materials have been deposited with the American Type Culture Collection, 10801 University Boulevard, Manassas, Virginia, USA (ATCC):

	<u>Material</u>	<u>ATCC Dep. No.</u>	<u>Deposit Date</u>
10	4E7.24.3	HB-12454	Jan. 13, 1998
	4H6.17.8	HB-12455	Jan. 13, 1998
	1H5.25.9	HB-12695	April 1, 1999
	4G7.18.8	PTA-99	May 21, 1999
	5G11.17.1	HB-12694	April 1, 1999
15	3F11.39.7	HB-12456	Jan. 13, 1998
	3H3.14.5	HB-12534	June 2, 1998

This deposit was made under the provisions of the Budapest Treaty on the International Recognition of the Deposit of Microorganisms for the Purpose of Patent Procedure and the Regulations thereunder (Budapest Treaty). This assures maintenance of a viable culture of the deposit for 30 years from the date of deposit. The deposit will be made available by ATCC under the terms of the Budapest Treaty, and subject to an agreement between Genentech, Inc. and ATCC, which assures permanent and unrestricted availability of the progeny of the culture of the deposit to the public upon issuance of the pertinent U.S. patent or upon laying open to the public of any U.S. or foreign patent application, whichever comes first, and assures availability of the progeny to one determined by the U.S. Commissioner of Patents and Trademarks to be entitled thereto according to 35 USC Section 122 and the Commissioner's rules pursuant thereto (including 37 CFR Section 1.14 with particular reference to 886 OG 638).

The assignee of the present application has agreed that if a culture of the materials on deposit should die or be lost or destroyed when cultivated under suitable conditions, the materials will be promptly replaced on notification with another of the same. Availability of the deposited material is not to be construed as a license to practice the invention in contravention of the rights granted under the authority of any government in accordance with its patent laws.

The foregoing written description is considered to be sufficient to enable one skilled in the art to practice the invention. The present invention is not to be limited in scope by the example presented herein. Indeed, various modifications of the invention in addition to those shown and described herein will become apparent to those skilled in the art from the foregoing description and fall within the scope of the appended claims.

## WHAT IS CLAIMED IS:

1. A method of enhancing apoptosis in mammalian cells, comprising exposing mammalian cells to an effective amount of CPT-11 and Apo-2 ligand receptor agonist, wherein said mammalian cells are exposed to the CPT-11 about 6 hours to about 72 hours prior to exposure to said Apo-2 ligand receptor agonist.
2. The method of claim 1 wherein the exposure of said mammalian cells to CPT-11 induces upregulation of DR4 receptor in said cells.
3. The method of claim 1 wherein the exposure of said mammalian cells to CPT-11 induces upregulation of DR5 receptor in said cells.
4. The method of claim 1 wherein said mammalian cells are exposed to CPT-11 about 24 or 48 hours prior to exposure to said Apo-2 ligand receptor agonist.
5. The method of claim 1 wherein said Apo-2 ligand receptor agonist comprises Apo2L polypeptide.
6. The method of claim 1 wherein said Apo-2 ligand receptor agonist comprises anti-DR4 receptor antibody.
7. The method of claim 6 wherein said anti-DR4 receptor antibody is a monoclonal antibody.
8. The method of claim 7 wherein said anti-DR4 receptor monoclonal antibody comprises a chimeric antibody.
9. The method of claim 7 wherein said anti-DR4 receptor monoclonal antibody comprises a human antibody.
10. The method of claim 1 wherein said Apo-2 ligand receptor agonist comprises anti-DR5 receptor antibody.
11. The method of claim 10 wherein said anti-DR5 receptor antibody is a monoclonal antibody.
12. The method of claim 11 wherein said anti-DR5 receptor monoclonal antibody comprises a chimeric antibody.
13. The method of claim 11 wherein said anti-DR5 receptor monoclonal antibody comprises a human antibody.
14. The method of claim 1 wherein said Apo-2 ligand receptor agonist is an anti-Apo-2 ligand receptor antibody which cross-reacts with more than one Apo-2 ligand receptor.
15. The method of claim 1 further comprising exposing the mammalian cells to one or more growth inhibitory agents.
16. The method of claim 1 further comprising exposing the mammalian cells to radiation.
17. The method of claim 1 wherein the mammalian cells are colorectal cancer cells.
18. A method of enhancing apoptosis in mammalian cancer cells, comprising exposing mammalian cells to an effective amount of CPT-11 and Apo-2 ligand receptor agonist, wherein (a) said mammalian cancer cells are exposed to the CPT-11 about 6 hours to about 72 hours prior

to exposure to said Apo-2 ligand receptor agonist and (b) said Apo-2 ligand receptor agonist is selected from the group consisting of Apo-2 ligand polypeptide comprising amino acid residues 114-281 of SEQ ID NO:1, anti-DR4 receptor antibody and anti-DR5 receptor antibody.

19. The method of claim 18 wherein the exposure of said mammalian cancer cells to CPT-11 induces upregulation of DR4 receptor in said cells.
20. The method of claim 18 wherein the exposure of said mammalian cancer cells to CPT-11 induces upregulation of DR5 receptor in said cells.
21. The method of claim 18 wherein said anti-DR4 receptor antibody or anti-DR5 receptor antibody is a chimeric, humanized or human antibody.
22. The method of claim 18 wherein said mammalian cancer cells are colorectal cancer cells.
23. The method of claim 18 wherein said Apo-2 ligand polypeptide consists of amino acid residues 114-281 of SEQ ID NO:1.
24. A method of treating cancer in a mammal, comprising administering to a mammal having cancer an effective amount of CPT-11 and Apo-2 ligand receptor agonist, wherein said CPT-11 is administered about 6 hours to about 72 hours prior to administration of the Apo-2 ligand receptor agonist.
25. The method of claim 24 wherein said Apo-2 ligand receptor agonist comprises Apo2L polypeptide.
26. The method of claim 24 wherein said Apo-2 ligand receptor agonist comprises an anti-DR4 receptor antibody.
27. The method of claim 24 wherein said Apo-2 ligand receptor agonist comprises an anti-DR5 receptor antibody.

1 / 14

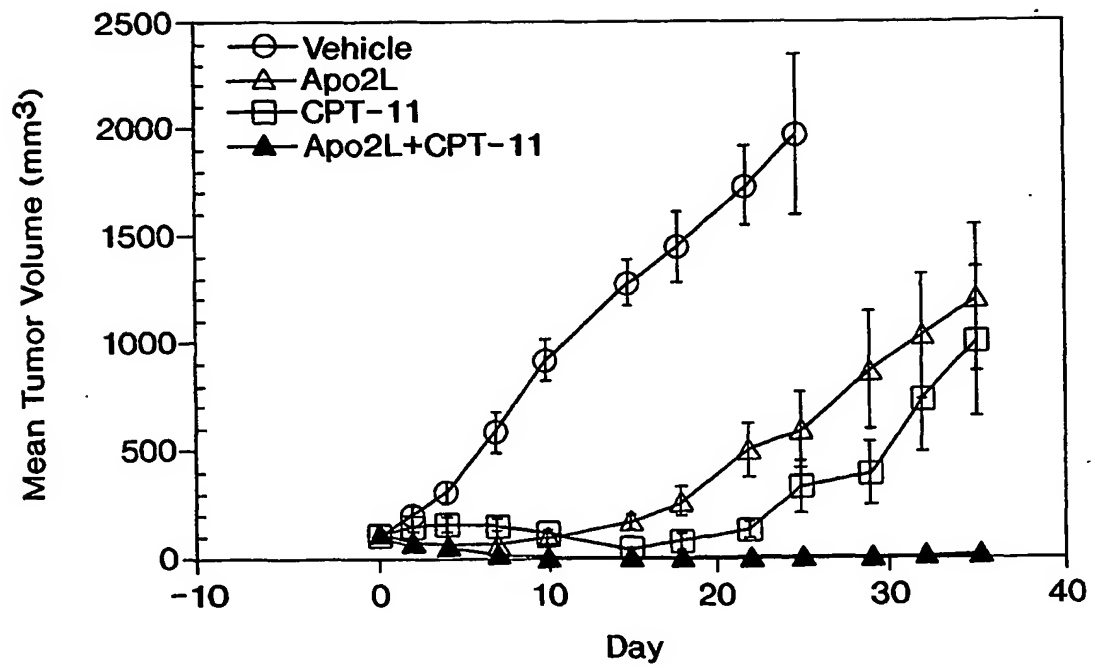


FIG. 1

2 / 14

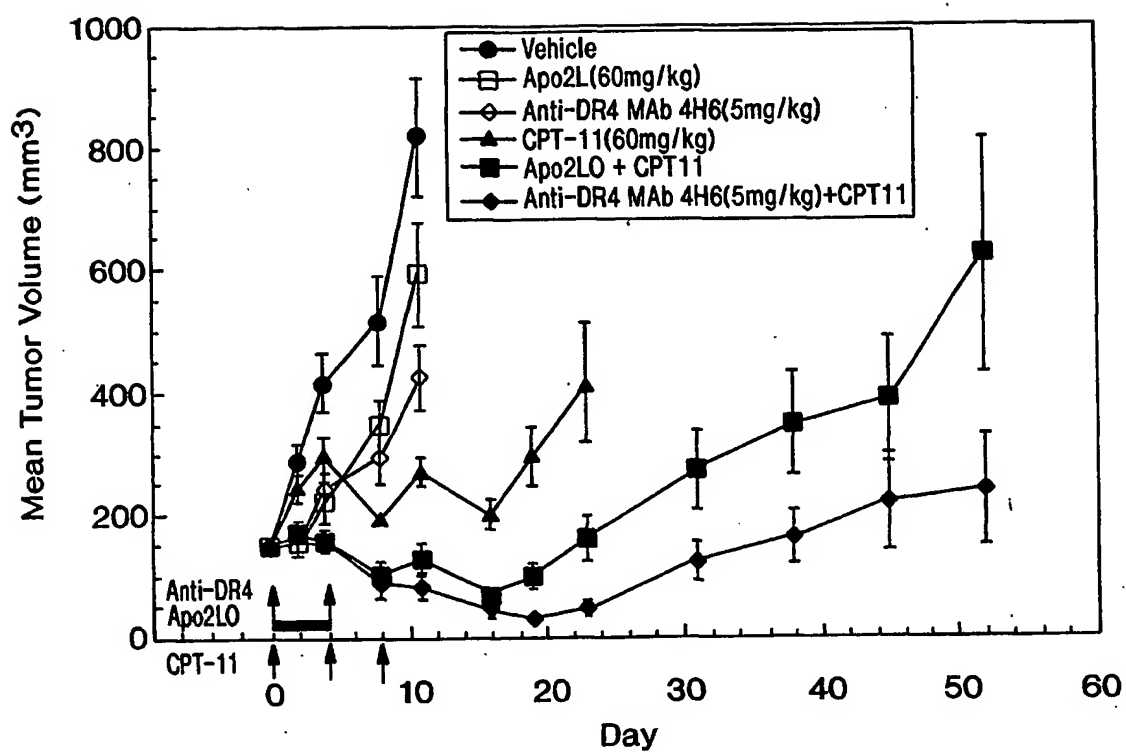
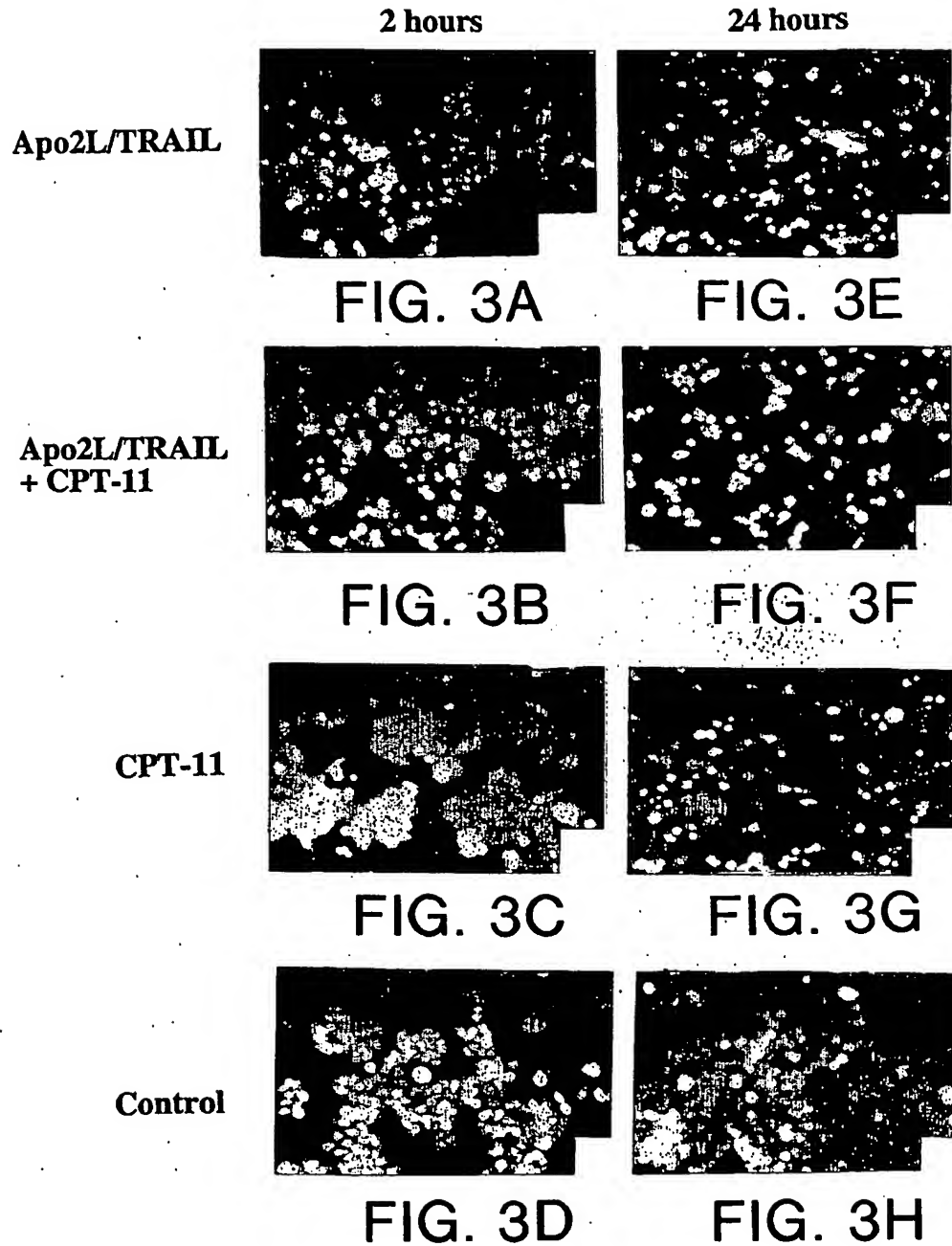


FIG. 2

3 / 14





4 / 14

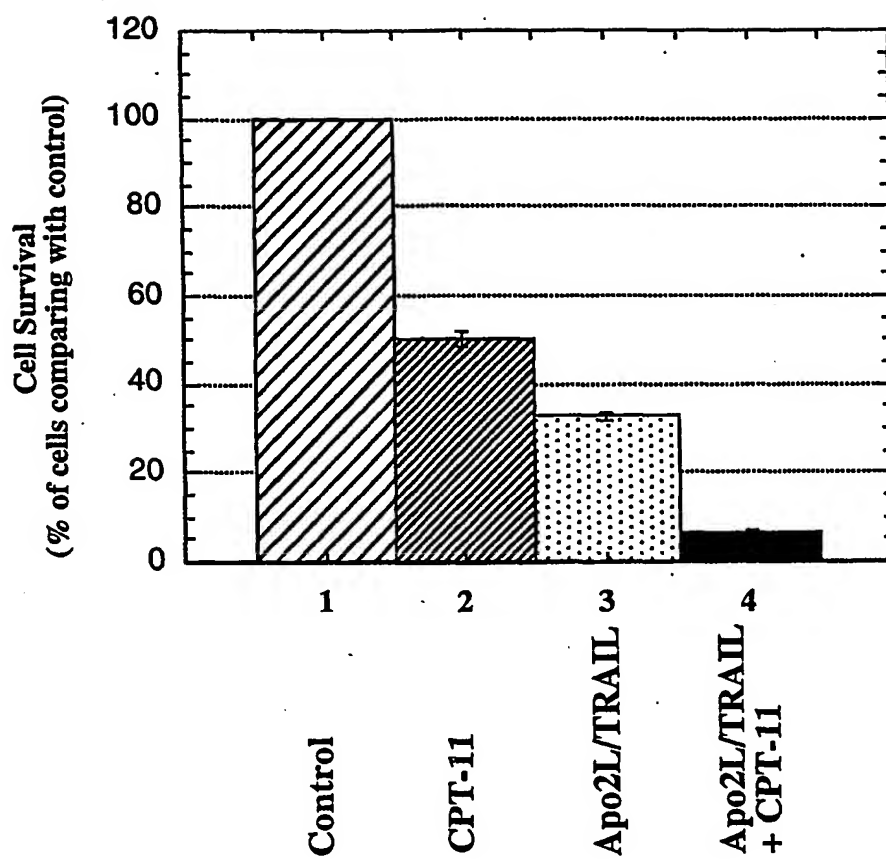


FIG. 4

5/14

FIG. 5A

FIG. 5B

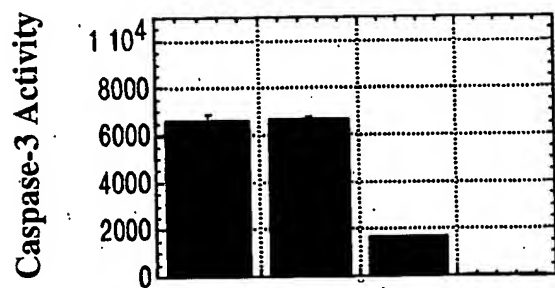
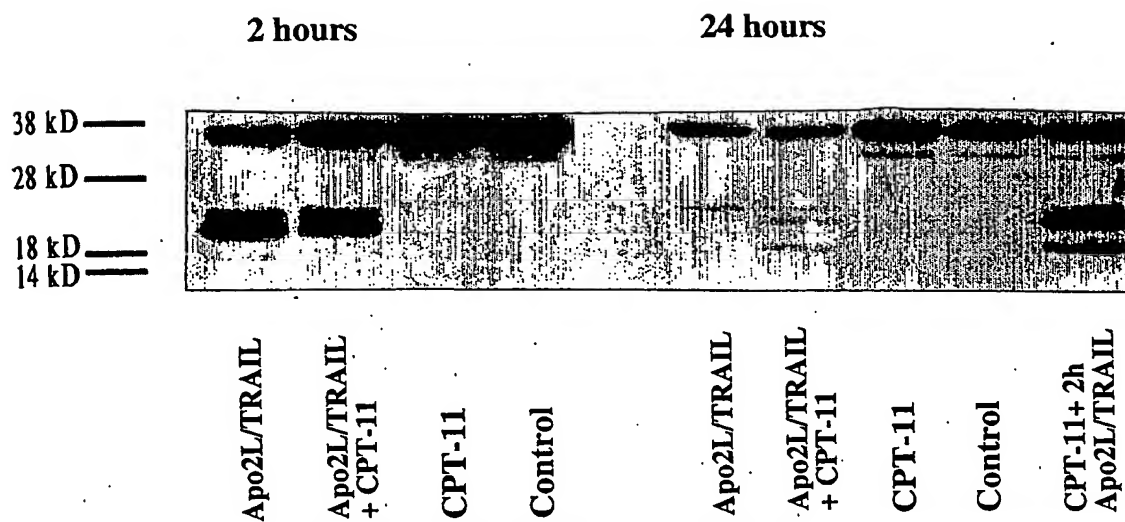


FIG. 5C

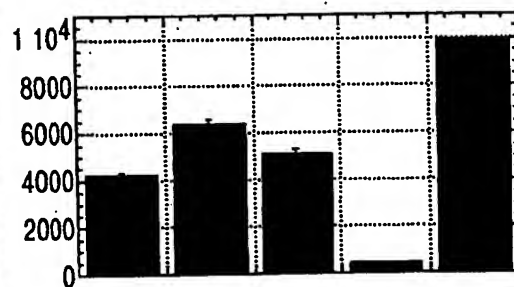


FIG. 5D

6 / 14

DR5

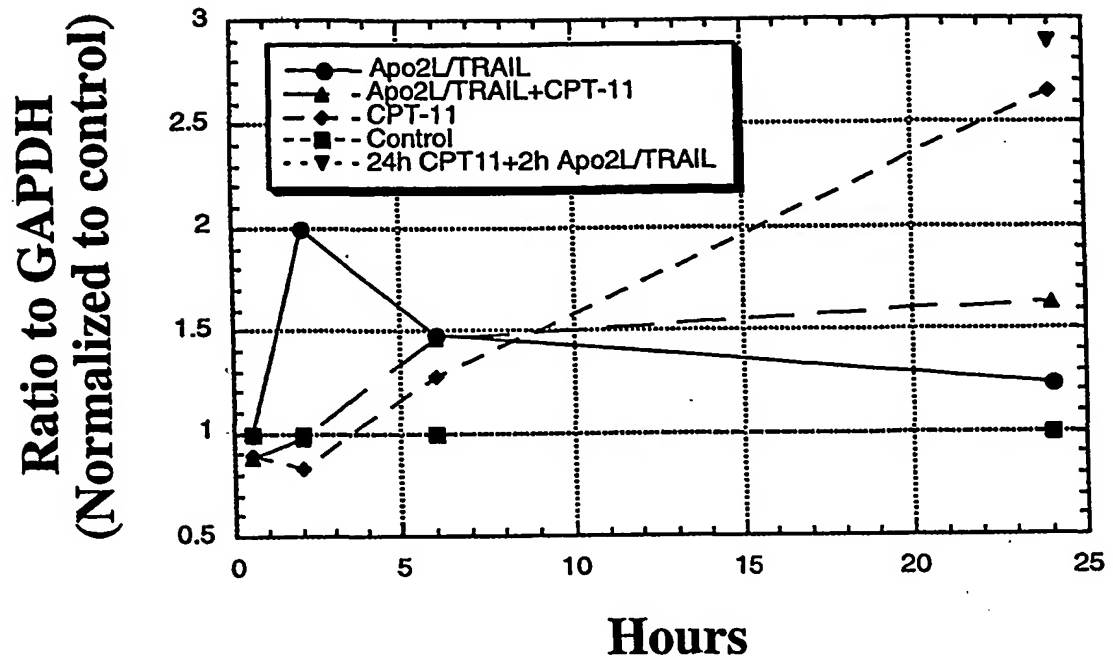


FIG. 6A

DR4

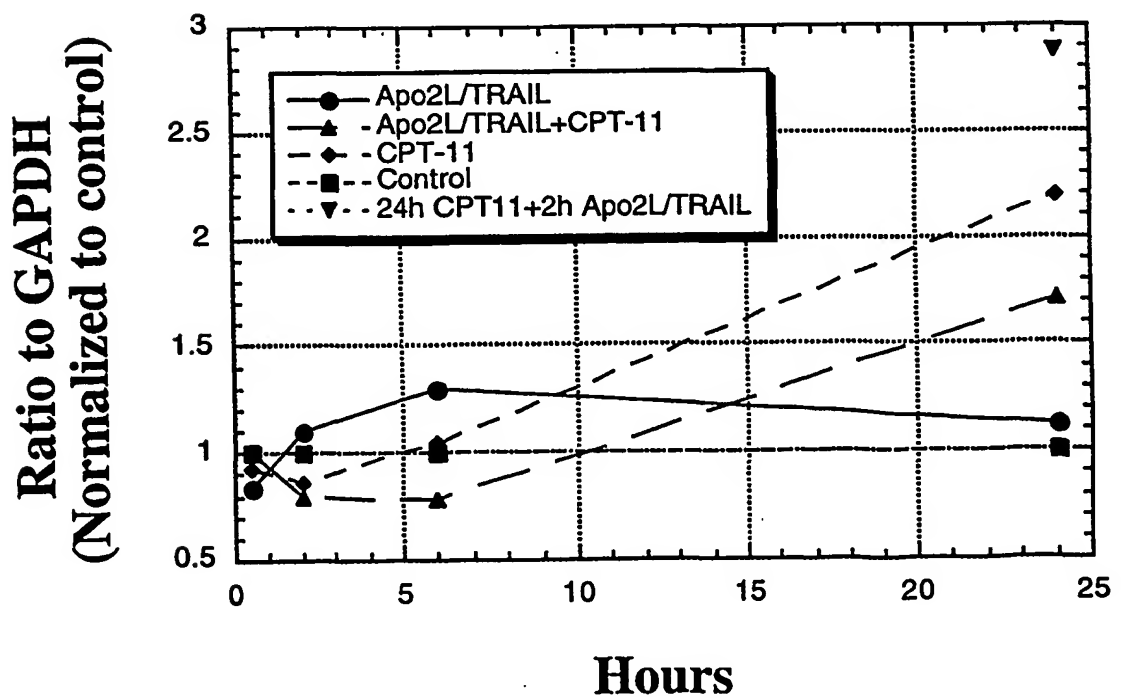


FIG. 6B

7/14

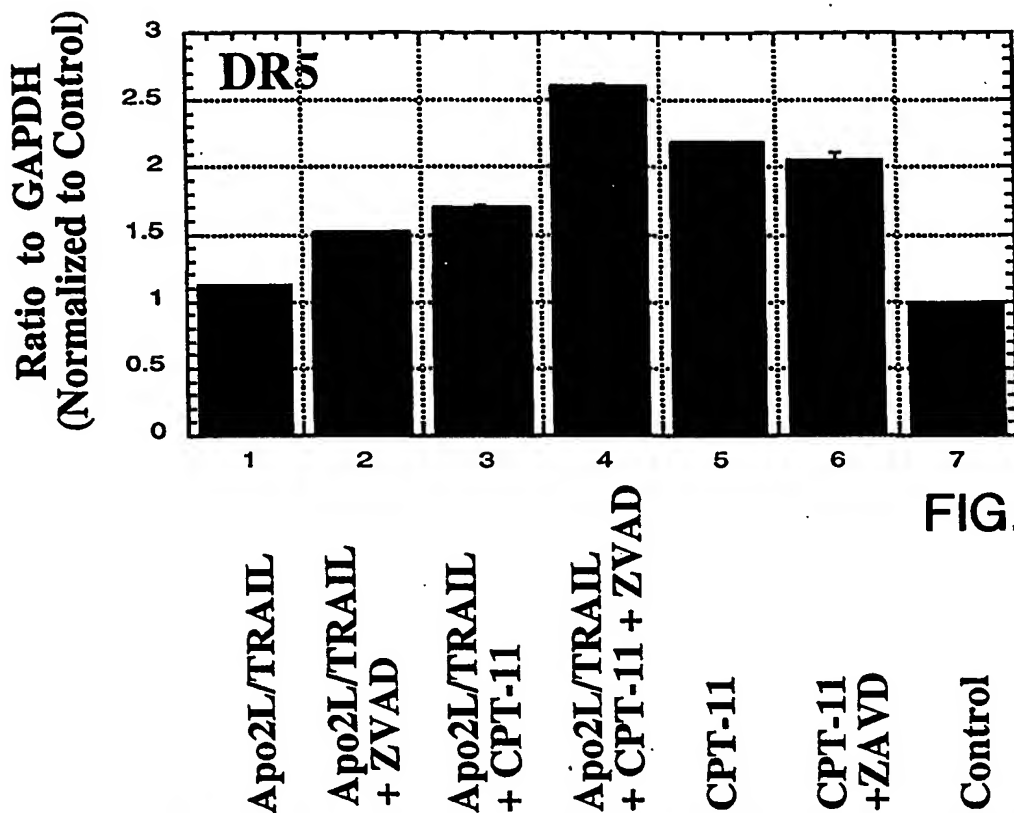


FIG. 7A

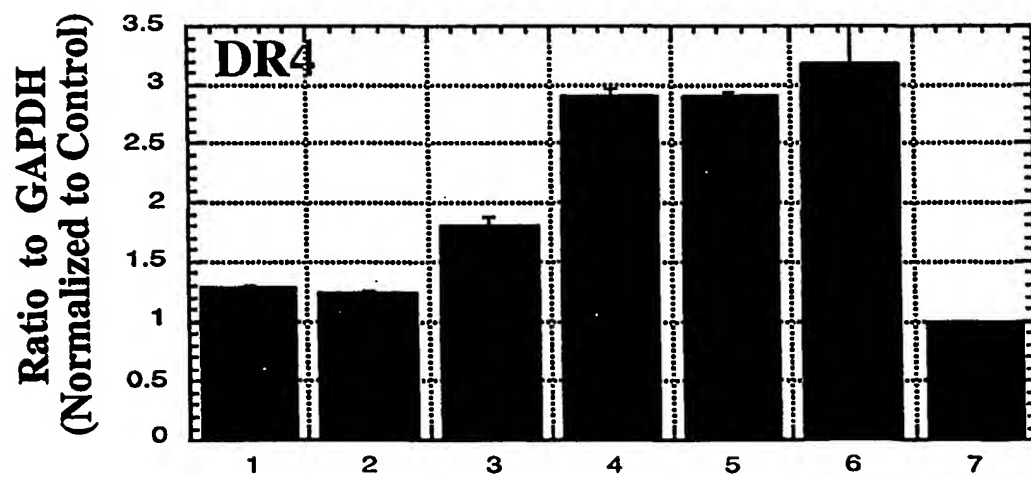


FIG. 7B

8/14

FIG. 8

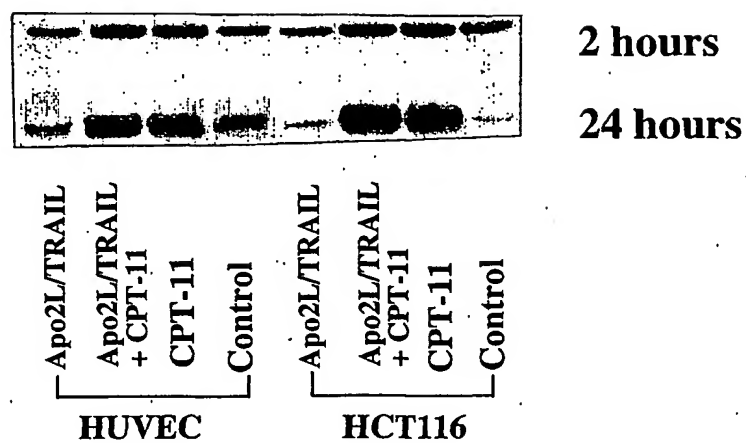
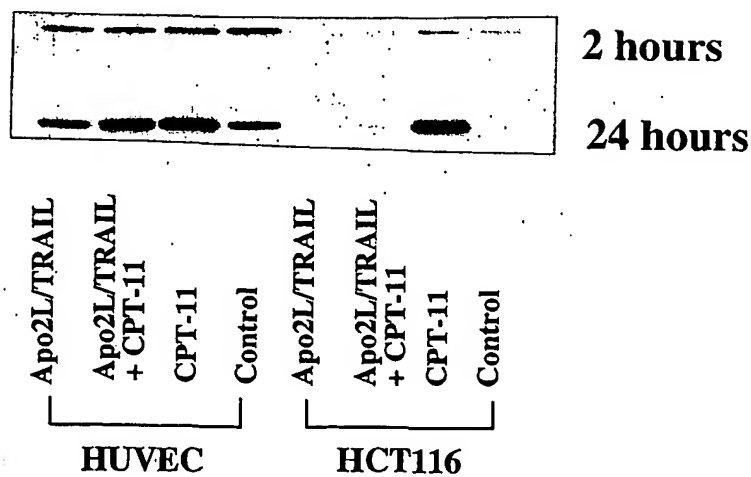
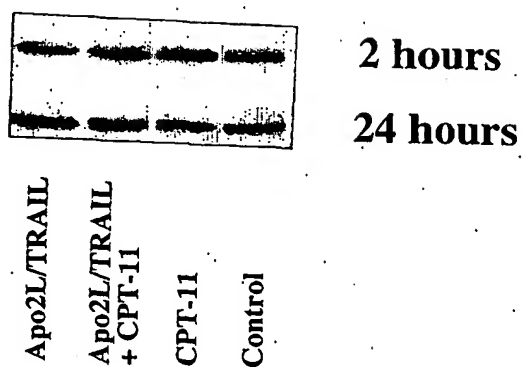


FIG. 9



9/14

FIG. 10



10/14

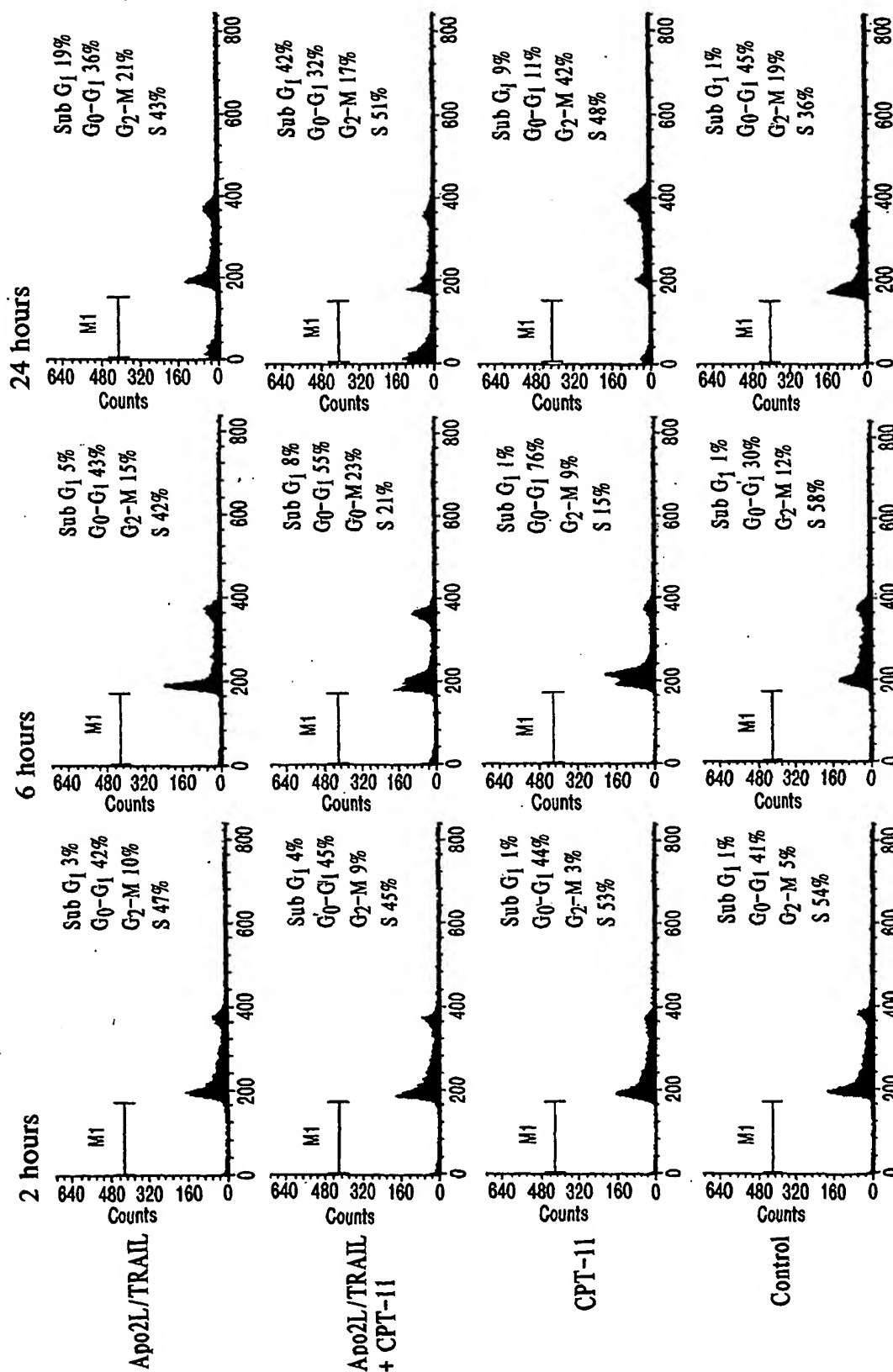
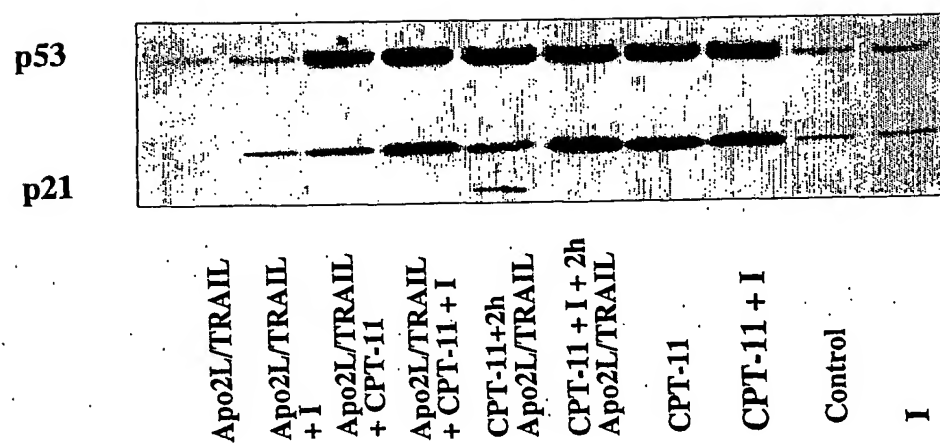


FIG. 11

11/14

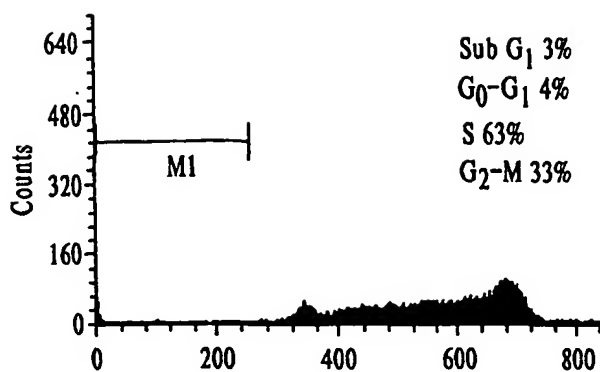
FIG. 12



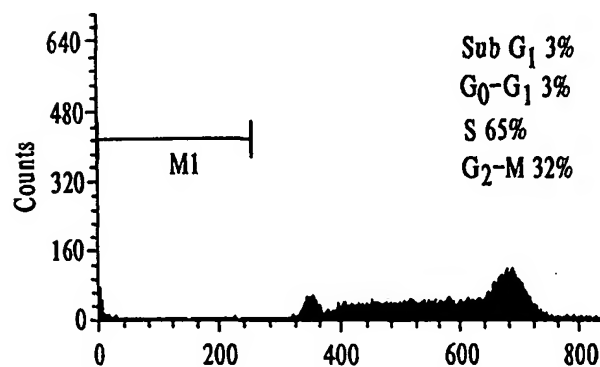


12/14

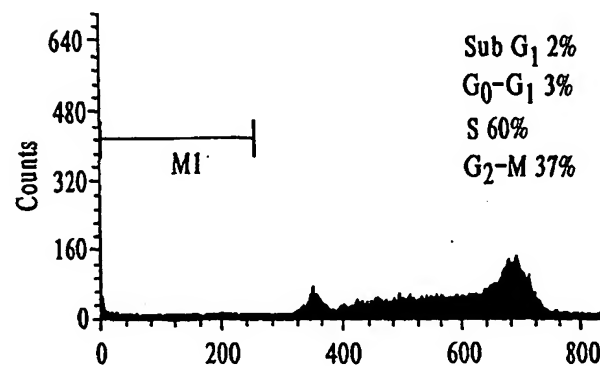
2h Apo2L/TRAIL  
+ CPT-11



Apo2L/TRAIL  
+ CPT-11



CPT-11



Control

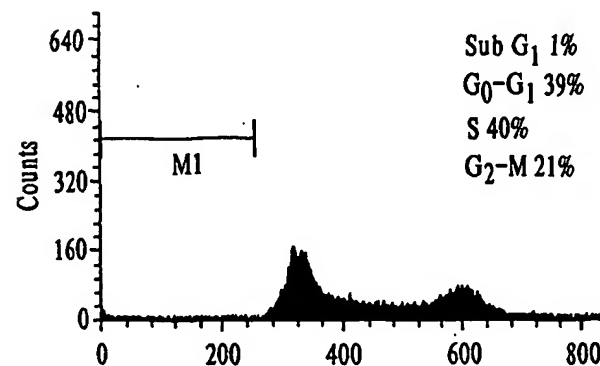


FIG. 13

13/14

FIG. 14A

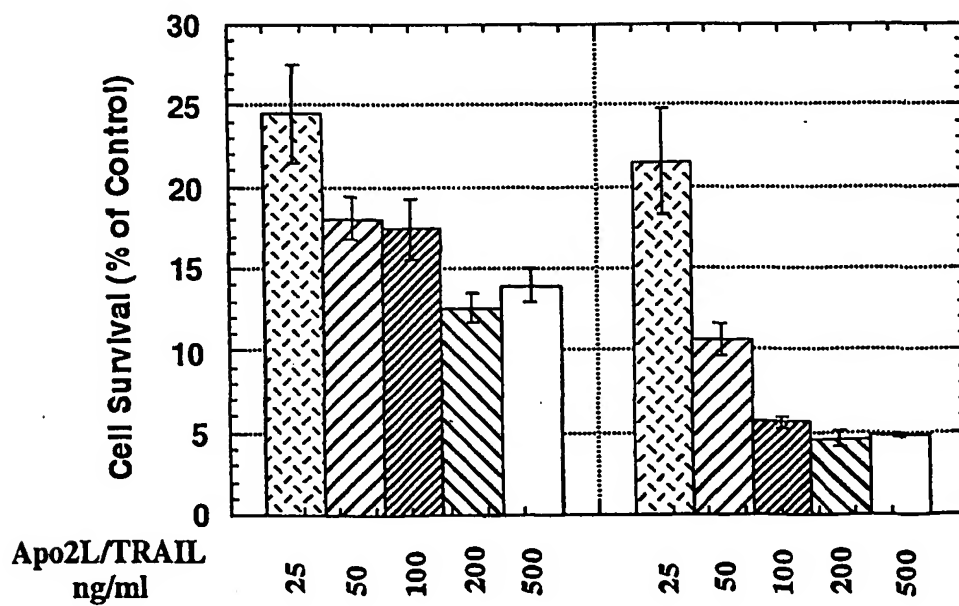
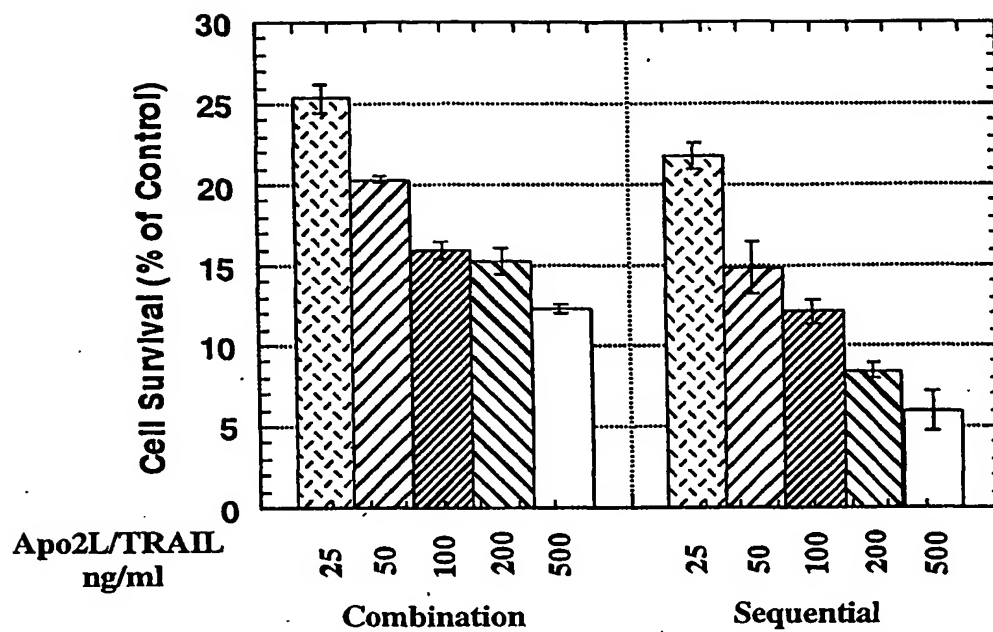


FIG. 14B

14/14

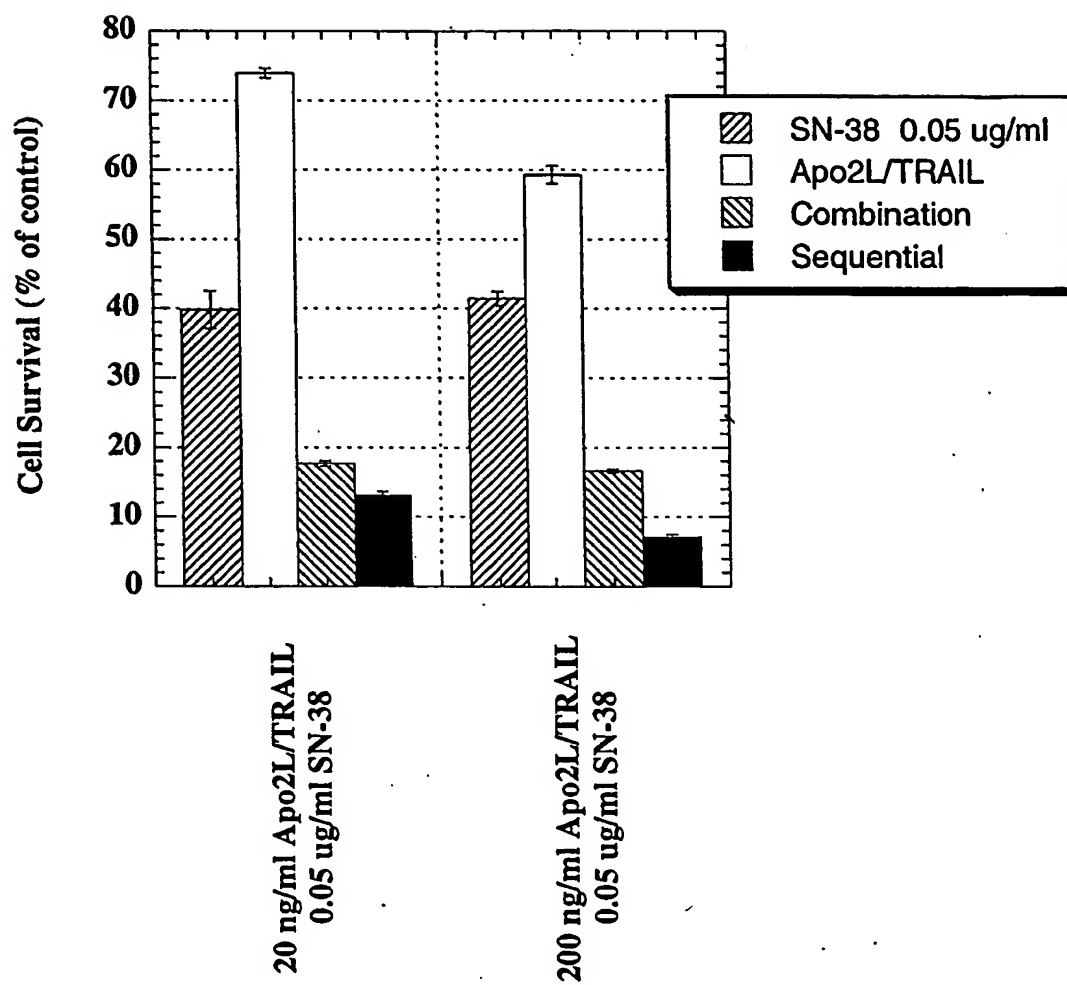


FIG. 15

## Sequence Listing

<110> GENENTECH, INC.  
 Escandon, Enrique  
 Fox, Judith A.  
 Kelley, Sean K.  
 Xiang, Hong

<120> APO-2L RECEPTOR AGONIST AND CPT-11 SYNERGISM

<130> P1838R1PCT

<141> 2001-07-27

<150> US 60/221,256  
 <151> 2000-07-27

<160> 4

<210> 1  
 <211> 281  
 <212> PRT  
 <213> Homo Sapien

<400> 1  
 Met Ala Met Met Glu Val Gln Gly Gly Pro Ser Leu Gly Gln Thr  
 1 5 10 15  
 Cys Val Leu Ile Val Ile Phe Thr Val Leu Leu Gln Ser Leu Cys  
 20 25 30  
 Val Ala Val Thr Tyr Val Tyr Phe Thr Asn Glu Leu Lys Gln Met  
 35 40 45  
 Gln Asp Lys Tyr Ser Lys Ser Gly Ile Ala Cys Phe Leu Lys Glu  
 50 55 60  
 Asp Asp Ser Tyr Trp Asp Pro Asn Asp Glu Glu Ser Met Asn Ser  
 65 70 75  
 Pro Cys Trp Gln Val Lys Trp Gln Leu Arg Gln Leu Val Arg Lys  
 80 85 90  
 Met Ile Leu Arg Thr Ser Glu Glu Thr Ile Ser Thr Val Gln Glu  
 95 100 105  
 Lys Gln Gln Asn Ile Ser Pro Leu Val Arg Glu Arg Gly Pro Gln  
 110 115 120  
 Arg Val Ala Ala His Ile Thr Gly Thr Arg Gly Arg Ser Asn Thr  
 125 130 135  
 Leu Ser Ser Pro Asn Ser Lys Asn Glu Lys Ala Leu Gly Arg Lys  
 140 145 150  
 Ile Asn Ser Trp Glu Ser Ser Arg Ser Gly His Ser Phe Leu Ser  
 155 160 165  
 Asn Leu His Leu Arg Asn Gly Glu Leu Val Ile His Glu Lys Gly  
 170 175 180  
 Phe Tyr Tyr Ile Tyr Ser Gln Thr Tyr Phe Arg Phe Gln Glu Glu  
 185 190 195  
 Ile Lys Glu Asn Thr Lys Asn Asp Lys Gln Met Val Gln Tyr Ile  
 200 205 210  
 Tyr Lys Tyr Thr Ser Tyr Pro Asp Pro Ile Leu Leu Met Lys Ser

215	220	225
Ala Arg Asn Ser Cys Trp Ser Lys Asp	Ala Glu Tyr Gly Leu Tyr	
230	235	240
Ser Ile Tyr Gln Gly Gly Ile Phe Glu	Leu Lys Glu Asn Asp Arg	
245	250	255
Ile Phe Val Ser Val Thr Asn Glu His	Leu Ile Asp Met Asp His	
260	265	270
Glu Ala Ser Phe Phe Gly Ala Phe Leu	Val Gly	
275	280	

<210> 2  
 <211> 468  
 <212> PRT  
 <213> Homo Sapien

<400> 2  
 Met Ala Pro Pro Pro Ala Arg Val His Leu Gly Ala Phe Leu Ala  
 1 5 10 15  
 Val Thr Pro Asn Pro Gly Ser Ala Ala Ser Gly Thr Glu Ala Ala  
 20 25 30  
 Ala Ala Thr Pro Ser Lys Val Trp Gly Ser Ser Ala Gly Arg Ile  
 35 40 45  
 Glu Pro Arg Gly Gly Gly Arg Gly Ala Leu Pro Thr Ser Met Gly  
 50 55 60  
 Gln His Gly Pro Ser Ala Arg Ala Arg Ala Gly Arg Ala Pro Gly  
 65 70 75  
 Pro Arg Pro Ala Arg Glu Ala Ser Pro Arg Leu Arg Val His Lys  
 80 85 90  
 Thr Phe Lys Phe Val Val Val Gly Val Leu Leu Gln Val Val Pro  
 95 100 105  
 Ser Ser Ala Ala Thr Ile Lys Leu His Asp Gln Ser Ile Gly Thr  
 110 115 120  
 Gln Gln Trp Glu His Ser Pro Leu Gly Glu Leu Cys Pro Pro Gly  
 125 130 135  
 Ser His Arg Ser Glu Arg Pro Gly Ala Cys Asn Arg Cys Thr Glu  
 140 145 150  
 Gly Val Gly Tyr Thr Asn Ala Ser Asn Asn Leu Phe Ala Cys Leu  
 155 160 165  
 Pro Cys Thr Ala Cys Lys Ser Asp Glu Glu Glu Arg Ser Pro Cys  
 170 175 180  
 Thr Thr Thr Arg Asn Thr Ala Cys Gln Cys Lys Pro Gly Thr Phe  
 185 190 195  
 Arg Asn Asp Asn Ser Ala Glu Met Cys Arg Lys Cys Ser Thr Gly  
 200 205 210  
 Cys Pro Arg Gly Met Val Lys Val Lys Asp Cys Thr Pro Trp Ser  
 215 220 225  
 Asp Ile Glu Cys Val His Lys Glu Ser Gly Asn Gly His Asn Ile  
 230 235 240

Trp Val Ile Leu Val Val Thr Leu Val Val Pro Leu Leu Leu Val  
 245 250 255  
 Ala Val Leu Ile Val Cys Cys Cys Ile Gly Ser Gly Cys Gly Gly  
 260 265 270  
 Asp Pro Lys Cys Met Asp Arg Val Cys Phe Trp Arg Leu Gly Leu  
 275 280 285  
 Leu Arg Gly Pro Gly Ala Glu Asp Asn Ala His Asn Glu Ile Leu  
 290 295 300  
 Ser Asn Ala Asp Ser Leu Ser Thr Phe Val Ser Glu Gln Gln Met  
 305 310 315  
 Glu Ser Gln Glu Pro Ala Asp Leu Thr Gly Val Thr Val Gln Ser  
 320 325 330  
 Pro Gly Glu Ala Gln Cys Leu Leu Gly Pro Ala Glu Ala Glu Gly  
 335 340 345  
 Ser Gln Arg Arg Arg Leu Leu Val Pro Ala Asn Gly Ala Asp Pro  
 350 355 360  
 Thr Glu Thr Leu Met Leu Phe Phe Asp Lys Phe Ala Asn Ile Val  
 365 370 375  
 Pro Phe Asp Ser Trp Asp Gln Leu Met Arg Gln Leu Asp Leu Thr  
 380 385 390  
 Lys Asn Glu Ile Asp Val Val Arg Ala Gly Thr Ala Gly Pro Gly  
 395 400 405  
 Asp Ala Leu Tyr Ala Met Leu Met Lys Trp Val Asn Lys Thr Gly  
 410 415 420  
 Arg Asn Ala Ser Ile His Thr Leu Leu Asp Ala Leu Glu Arg Met  
 425 430 435  
 Glu Glu Arg His Ala Lys Glu Lys Ile Gln Asp Leu Leu Val Asp  
 440 445 450  
 Ser Gly Lys Phe Ile Tyr Leu Glu Asp Gly Thr Gly Ser Ala Val  
 455 460 465  
 Ser Leu Glu

&lt;210&gt; 3

&lt;211&gt; 440

&lt;212&gt; PRT

&lt;213&gt; Homo Sapien

&lt;400&gt; 3

Met Glu Gln Arg Gly Gln Asn Ala Pro Ala Ala Ser Gly Ala Arg  
 1 5 10 15  
 Lys Arg His Gly Pro Gly Pro Arg Glu Ala Arg Gly Ala Arg Pro  
 20 25 30  
 Gly Pro Arg Val Pro Lys Thr Leu Val Leu Val Val Ala Ala Val  
 35 40 45  
 Leu Leu Leu Val Ser Ala Glu Ser Ala Leu Ile Thr Gln Gln Asp  
 50 55 60  
 Leu Ala Pro Gln Gln Arg Ala Ala Pro Gln Gln Lys Arg Ser Ser  
 65 70 75

Pro	Ser	Glu	Gly	Leu	Cys	Pro	Pro	Gly	His	His	Ile	Ser	Glu	Asp	
				80					85					90	
Gly	Arg	Asp	Cys	Ile	Ser	Cys	Lys	Tyr	Gly	Gln	Asp	Tyr	Ser	Thr	
				95					100					105	
His	Trp	Asn	Asp	Leu	Leu	Phe	Cys	Leu	Arg	Cys	Thr	Arg	Cys	Asp	
				110					115					120	
Ser	Gly	Glu	Val	Glu	Leu	Ser	Pro	Cys	Thr	Thr	Thr	Arg	Asn	Thr	
				125					130					135	
Val	Cys	Gln	Cys	Glu	Glu	Gly	Thr	Phe	Arg	Glu	Glu	Asp	Ser	Pro	
				140					145					150	
Glu	Met	Cys	Arg	Lys	Cys	Arg	Thr	Gly	Cys	Pro	Arg	Gly	Met	Val	
				155					160					165	
Lys	Val	Gly	Asp	Cys	Thr	Pro	Trp	Ser	Asp	Ile	Glu	Cys	Val	His	
				170					175					180	
Lys	Glu	Ser	Gly	Thr	Lys	His	Ser	Gly	Glu	Ala	Pro	Ala	Val	Glu	
				185					190					195	
Glu	Thr	Val	Thr	Ser	Ser	Pro	Gly	Thr	Pro	Ala	Ser	Pro	Cys	Ser	
				200					205					210	
Leu	Ser	Gly	Ile	Ile	Ile	Gly	Val	Thr	Val	Ala	Ala	Val	Val	Leu	
				215					220					225	
Ile	Val	Ala	Val	Phe	Val	Cys	Lys	Ser	Leu	Leu	Trp	Lys	Lys	Val	
				230					235					240	
Leu	Pro	Tyr	Leu	Lys	Gly	Ile	Cys	Ser	Gly	Gly	Gly	Gly	Asp	Pro	
				245					250					255	
Glu	Arg	Val	Asp	Arg	Ser	Ser	Gln	Arg	Pro	Gly	Ala	Glu	Asp	Asn	
				260					265					270	
Val	Leu	Asn	Glu	Ile	Val	Ser	Ile	Leu	Gln	Pro	Thr	Gln	Val	Pro	
				275					280					285	
Glu	Gln	Glu	Met	Glu	Val	Gln	Glu	Pro	Ala	Glu	Pro	Thr	Gly	Val	
				290					295					300	
Asn	Met	Leu	Ser	Pro	Gly	Glu	Ser	Glu	His	Leu	Leu	Glu	Pro	Ala	
				305					310					315	
Glu	Ala	Glu	Arg	Ser	Gln	Arg	Arg	Arg	Leu	Leu	Val	Pro	Ala	Asn	
				320					325					330	
Glu	Gly	Asp	Pro	Thr	Glu	Thr	Leu	Arg	Gln	Cys	Phe	Asp	Asp	Phe	
				335					340					345	
Ala	Asp	Leu	Val	Pro	Phe	Asp	Ser	Trp	Glu	Pro	Leu	Met	Arg	Lys	
				350					355					360	
Leu	Gly	Leu	Met	Asp	Asn	Glu	Ile	Lys	Val	Ala	Lys	Ala	Glu	Ala	
				365					370					375	
Ala	Gly	His	Arg	Asp	Thr	Leu	Tyr	Thr	Met	Leu	Ile	Lys	Trp	Val	
				380					385					390	
Asn	Lys	Thr	Gly	Arg	Asp	Ala	Ser	Val	His	Thr	Leu	Leu	Asp	Ala	
				395					400					405	
Leu	Glu	Thr	Leu	Gly	Glu	Arg	Leu	Ala	Lys	Gln	Lys	Ile	Glu	Asp	

410	415	420
His Leu Leu Ser Ser Gly Lys Phe Met Tyr Leu Glu Gly Asn Ala		
425	430	435
Asp Ser Ala Met Ser		
440		
<210> 4		
<211> 411		
<212> PRT		
<213> Homo Sapien		
<220>		
<221> unsure		
<222> 410		
<223> unknown amino acid		
<400> 4		
Met Glu Gln Arg Gly Gln Asn Ala Pro Ala Ala Ser Gly Ala Arg		
1 5 10 15		
Lys Arg His Gly Pro Gly Pro Arg Glu Ala Arg Gly Ala Arg Pro		
20 25 30		
Gly Leu Arg Val Pro Lys Thr Leu Val Leu Val Val Ala Ala Val		
35 40 45		
Leu Leu Leu Val Ser Ala Glu Ser Ala Leu Ile Thr Gln Gln Asp		
50 55 60		
Leu Ala Pro Gln Gln Arg Ala Ala Pro Gln Gln Lys Arg Ser Ser		
65 70 75		
Pro Ser Glu Gly Leu Cys Pro Pro Gly His His Ile Ser Glu Asp		
80 85 90		
Gly Arg Asp Cys Ile Ser Cys Lys Tyr Gly Gln Asp Tyr Ser Thr		
95 100 105		
His Trp Asn Asp Leu Leu Phe Cys Leu Arg Cys Thr Arg Cys Asp		
110 115 120		
Ser Gly Glu Val Glu Leu Ser Pro Cys Thr Thr Thr Arg Asn Thr		
125 130 135		
Val Cys Gln Cys Glu Glu Gly Thr Phe Arg Glu Glu Asp Ser Pro		
140 145 150		
Glu Met Cys Arg Lys Cys Arg Thr Gly Cys Pro Arg Gly Met Val		
155 160 165		
Lys Val Gly Asp Cys Thr Pro Trp Ser Asp Ile Glu Cys Val His		
170 175 180		
Lys Glu Ser Gly Ile Ile Ile Gly Val Thr Val Ala Ala Val Val		
185 190 195		
Leu Ile Val Ala Val Phe Val Cys Lys Ser Leu Leu Trp Lys Lys		
200 205 210		
Val Leu Pro Tyr Leu Lys Gly Ile Cys Ser Gly Gly Gly Gly Asp		
215 220 225		
Pro Glu Arg Val Asp Arg Ser Ser Gln Arg Pro Gly Ala Glu Asp		
230 235 240		
Asn Val Leu Asn Glu Ile Val Ser Ile Leu Gln Pro Thr Gln Val		



245	250	255
Pro Glu Gln Glu Met Glu Val Gln Glu	Pro Ala Glu Pro Thr Gly	
260	265	270
Val Asn Met Leu Ser Pro Gly Glu Ser	Glu His Leu Leu Glu Pro	
275	280	285
Ala Glu Ala Glu Arg Ser Gln Arg Arg	Arg Leu Leu Val Pro Ala	
290	295	300
Asn Glu Gly Asp Pro Thr Glu Thr Leu	Arg Gln Cys Phe Asp Asp	
305	310	315
Phe Ala Asp Leu Val Pro Phe Asp Ser	Trp Glu Pro Leu Met Arg	
320	325	330
Lys Leu Gly Leu Met Asp Asn Glu Ile	Lys Val Ala Lys Ala Glu	
335	340	345
Ala Ala Gly His Arg Asp Thr Leu Tyr	Thr Met Leu Ile Lys Trp	
350	355	360
Val Asn Lys Thr Gly Arg Asp Ala Ser	Val His Thr Leu Leu Asp	
365	370	375
Ala Leu Glu Thr Leu Gly Glu Arg Leu	Ala Lys Gln Lys Ile Glu	
380	385	390
Asp His Leu Leu Ser Ser Gly Lys Phe	Met Tyr Leu Glu Gly Asn	
395	400	405
Ala Asp Ser Ala Xaa Ser		
410		

(19) World Intellectual Property Organization  
International Bureau



(43) International Publication Date  
7 February 2002 (07.02.2002)

PCT

(10) International Publication Number  
**WO 02/009755 A3**

(51) International Patent Classification<sup>7</sup>: **A61K 38/17**,  
39/395, 31/47, A61P 35/00, 37/00 // (A61K 39/395,  
31:47) (A61K 38/17, 31:47)

(21) International Application Number: PCT/US01/23691

(22) International Filing Date: 27 July 2001 (27.07.2001)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:  
60/221,256 27 July 2000 (27.07.2000) US

(71) Applicant (for all designated States except US): GENEN-  
TECH, INC. [US/US]; 1 DNA Way, South San Francisco,  
CA 94080 (US).

(72) Inventors; and

(75) Inventors/Applicants (for US only): ESCANDON, En-  
rique [MX/US]; 10 Garden Ct. Apt. #2, Belmont, CA  
94002 (US). FOX, Judith, A. [US/US]; 1531 Golden Gate  
Avenue, San Francisco, CA 94115 (US). KELLEY, Sean,  
K. [US/US]; 3915 Edison Street, San Mateo, CA 94403  
(US). XIANG, Hong [US/US]; 558 Driscoll Place, Palo  
Alto, CA 94306 (US).

(74) Agent: MARSCHANG, Diane, L.; Genentech, Inc., MS  
49, 1 DNA Way, South San Francisco, CA 94080 (US).

(81) Designated States (national): AE, AG, AL, AM, AT, AU,  
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,  
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,  
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,  
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,  
MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,  
SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA,  
ZW.

(84) Designated States (regional): ARIPO patent (GH, GM,  
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian  
patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European  
patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE,  
IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF,  
CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD,  
TG).

**Published:**

- with international search report
- with sequence listing part of description published sepa-  
rately in electronic form and available upon request from  
the International Bureau

(88) Date of publication of the international search report:  
13 February 2003

For two-letter codes and other abbreviations, refer to the "Guid-  
ance Notes on Codes and Abbreviations" appearing at the begin-  
ning of each regular issue of the PCT Gazette.



**WO 02/009755 A3**

(54) Title: APO-2L RECEPTOR AGONIST AND CPT-11 SYNERGISM

(57) Abstract: Methods of using effective amounts of Apo-2L receptor agonists and CPT-11 to induce apoptosis and suppress growth of cancer cells are provided.

## INTERNATIONAL SEARCH REPORT

Int. Appl. No.  
PCT/US 01/23691

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 A61K38/17 A61K39/395 A61K31/47 A61P35/00 A61P37/00  
 //(A61K39/395,31:47),(A61K38/17,31:47)

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the International search (name of data base and, where practical, search terms used)

BIOSIS, EMBASE, WPI Data, PAJ, EPO-Internal, SEQUENCE SEARCH

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	A. ASHKENAZI ET AL.: "Safety and antitumor activity of recombinant soluble Apo2 ligand." JOURNAL OF CLINICAL INVESTIGATION, vol. 104, no. 2, July 1999 (1999-07), pages 155-162, XP000973369 New York, NY, USA abstract page 161, left-hand column, line 2 - line 16 page 162, left-hand column, line 6 - line 29	1-5, 15, 17-20, 22-25
Y	---	6-13, 16, 21, 26, 27
	--- -/--	

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

## \* Special categories of cited documents:

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

- \*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- \*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- \*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
- \*&\* document member of the same patent family

Date of the actual completion of the international search

23 July 2002

Date of mailing of the international search report

02/08/2002

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2  
 NL - 2280 HV Rijswijk  
 Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,  
 Fax: (+31-70) 340-3016

Authorized officer

No01J, F

Internal Application No  
PCR/US 01/23691

Form PCT/ISA/210 (continuation of second sheet) (July 1992)

## INTERNATIONAL SEARCH REPORT

Int. Patent Application No.  
PCT/US 01/23691

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P, X	WO 00 75191 A (GENENTECH INC.) 14 December 2000 (2000-12-14) examples claims	1-27
P, X	H. XIANG ET AL.: "Combined Apo2L/TRAIL and CPT-11 enhanced tumor cell apoptosis." PROCEEDINGS OF THE AMERICAN ASSOCIATION FOR CANCER RESEARCH, vol. 42, March 2001 (2001-03), page 643 XP002207072 Philadelphia, PA, USA abstract #3457	1-5, 17-20, 22-25

# INTERNATIONAL SEARCH REPORT

ational application No.  
PCT/US 01/23691

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
Although claims 1-23 (all partially, as far as an in vivo method is concerned) and claims 24-27 (all completely) are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this International application, as follows:

see additional sheet

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☒ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: 5, 23, 25 (all completely), 1-4, 15-20, 22, 24 (all partially)

Methods of using effective amounts of Apo-2L receptor agonists and CPT-11 to induce apoptosis and suppress growth of mammalian cancer cells, wherein said Apo-2L receptor agonist comprises Apo-2L polypeptide.

2. Claims: 6-9, 26 (all completely), 1-4, 14-20, 21, 22, 24 (all partially)

Methods of using effective amounts of Apo-2L receptor agonists and CPT-11 to induce apoptosis and suppress growth of mammalian cancer cells, wherein said Apo-2L receptor agonists comprises anti-DR4 receptor antibody.

3. Claims: 10-13, 27 (all completely), 1-4, 14-20, 21, 22, 24 (all partially)

Methods of using effective amounts of Apo-2L receptor agonists and CPT-11 to induce apoptosis and suppress growth of mammalian cancer cells, wherein said Apo-2L receptor agonists comprises anti-DR5 receptor antibody.

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No  
PCT/JS 01/23691

Patent document cited in search report		Publication date	Patent family member(s)	Publication date
WO 9948527	A	30-09-1999	AU 3365999 A CA 2324494 A1 EP 1064027 A1 JP 2002507404 T WO 9948527 A1	18-10-1999 30-09-1999 03-01-2001 12-03-2002 30-09-1999
WO 9937684	A	29-07-1999	AU 2338299 A CA 2318405 A1 EP 1053256 A1 JP 2002500877 T WO 9937684 A1	09-08-1999 29-07-1999 22-11-2000 15-01-2002 29-07-1999
WO 0075191	A	14-12-2000	AU 5596500 A EP 1192185 A2 WO 0075191 A2	28-12-2000 03-04-2002 14-12-2000